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Thyroid hormone modulation during zebrafish development recapitulates evolved diversity in danionin jaw protrusion mechanics

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Abstract

Protrusile jaws are a highly useful innovation that has been linked to extensive diversification in fish feeding ecology. Jaw protrusion can enhance the performance of multiple functions, such as suction production and capturing elusive prey. Identifying the developmental factors that alter protrusion ability will improve our understanding of fish diversification. In the zebrafish protrusion arises postmetamorphosis. Fish metamorphosis typically includes significant changes in trophic morphology, accompanies a shift in feeding niche and coincides with increased thyroid hormone production. We tested whether thyroid hormone affects the development of zebrafish feeding mechanics. We found that it affected all developmental stages examined, but that effects were most pronounced after metamorphosis. Thyroid hormone levels affected the development of jaw morphology, feeding mechanics, shape variation, and cranial ossification. Adult zebrafish utilize protrusile jaws, but an absence of thyroid hormone impaired development of the premaxillary bone, which is critical to jaw protrusion. Premaxillae from early juvenile zebrafish and hypothyroid adult zebrafish resemble those from adults in the genera *Danionella*, *Devario*, and *Microdevario* that show little to no jaw protrusion. Our findings suggest that evolutionary changes in how the developing skulls of danionin minnows respond to thyroid hormone may have promoted diversification into different feeding niches.

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AUTHOR CONTRIBUTIONS

D. G. conducted this research as the subject of her master's thesis, performed experiments, collected data, analysed data and contributed to the writing the manuscript; E. S. oversaw graduate and undergraduate student training in the techniques used during this research, coordinated student research efforts and provided comments on the manuscript; Z. D. collected and analysed data; M. W. collected and analysed data; A.D. collected and analysed data; C. C. collected and analysed data; S. K. M. analysed data and provided comments on the manuscript; W. J. C. oversaw the research project, analysed data, and contributed to the writing the manuscript. W. J. C. designed and performed the study, analyzed data, prepared and wrote the manuscript. D. G., E. S., Z. D., M. W., A. D. and C. C. performed experiments and analyzed data. S. K. M. provided technical advice and revised the manuscript.

SUPPORTING INFORMATION

Additional supporting information may be found online in the Supporting Information section.

CONFLICT OF INTERESTS

The authors declare that there are no conflict of interests.

Keywords

functional morphology of feeding; metamorphosis; skull

1 | INTRODUCTION

Many if not most fishes undergo a metamorphosis during which their bodies are extensively remodeled (McMenamin & Parichy, 2013). It is during metamorphosis and postmetamorphic development that most fishes acquire the behavioral and morphological characters that allow them to occupy their adult niches. Premetamorphic (larval) and postmetamorphic (juvenile and adult) fishes of the same species frequently live in different habitats and occupy disparate feeding niches (Leis & McCormick, 2006; McCormick & Makey, 1997; McCormick, Makey, & Dufour, 2002; McMenamin & Parichy, 2013). The ecological diversification of adults in many fish lineages is therefore closely linked to evolutionary changes in the processes that shape their postlarval development. Most molecular developmental studies have focused on early development, particularly embryonic stages (Albertson & Yelick, 2004; Cooper, Wirgau, Sweet, & Albertson, 2013; McMenamin & Parichy, 2013; Parsons, Andreeva, Cooper, Yelick, & Albertson, 2010), but if we are to understand the developmental changes that have permitted the adaptive diversification of adult fish feeding we need a better understanding of the controls of morphogenesis in late development.

Protrusile jaws are an important evolutionary innovation in fish feeding. Highly moveable skull linkages that allow the jaws to protrude forward from the face have evolved independently in at least six lineages of bony fishes and protrusion has been lost, gained, reduced, and enhanced many times in these clades (Ferry-Graham, Gibb, & Hernandez, 2008; Staab, Holzman, Hernandez, & Wainwright, 2012; Wainwright, McGee, Longo, & Hernandez, 2015). Two of these, Cypriniformes (~3,200 species, including the zebrafish) and Acanthomorpha (~17,000 species), have been particularly successful and together comprise more than one-third of living vertebrates (Near et al., 2013; Staab et al., 2012; Wainwright et al., 2015; Yang et al., 2010). Maximum jaw protrusion distance has been closely linked with diet (Cooper, Carter, Conith, Rice, & Westneat, 2017) and an ability to rapidly transition between morphs capable of different degrees of protrusion appears to support diversification into different feeding niches (Cooper & Westneat, 2009; Cooper et al., 2010; Ferry-Graham et al., 2008; Staab et al., 2012; Wainwright et al., 2015).

Although protrusile jaws can confer multiple functional abilities in adults, particularly the enhancement of suction production via rapid expansion of the mouth cavity (i.e., buccal cavity; Ferry-Graham et al., 2008; Holzman, Collar, Mehta, & Wainwright, 2012; Konow & Bellwood, 2005; Staab et al., 2012), the relatively high water viscosities experienced by small aquatic organisms severely limits the utility of protrusile jaws in fish larvae (Hernandez, 2000; Hernández, Barresi, & Devoto, 2002; Yaniv, Elad, & Holzman, 2014). Small fishes live in a low Reynolds number environment in which viscous forces are greater than inertial forces (Hernandez, 1995; Hernandez, 2000). These animals experience water as a “sticky” fluid that can reduce the effectiveness of feeding techniques that are viable at

larger body sizes (Hernandez, 1995; Hernandez, 2000). Protrusile jaws would most likely interfere with suction feeding in fish larvae and jaw protrusion does not arise until after metamorphosis in the zebrafish (Hernández et al., 2002; McMenamin, Carter, & Cooper, 2017).

The reshaping of multiple skull bones must be coordinated during the larva-to-juvenile transition for a fish to have a properly integrated mature skull. This need for integrated development should be particularly strong for fishes with complex, highly kinetic adult skulls in which motion is transferred through multiple linkages. Hormones reach all body organs essentially simultaneously via circulating blood, and therefore have the potential to act as agents of developmental coordination that stimulate multiple organs, including different bones, to transform at the same time. Along with additional endocrine factors, thyroid hormone (TH) signaling plays a major role in directing late-developmental skeletal remodeling in vertebrates, and TH stimulates metamorphosis or metamorphosis-like processes in many species (Das et al., 2006; Hu et al., 2019; Laudet, 2011; McMenamin & Parichy, 2013; Paris et al., 2010; Wojcicka, Bassett, & Williams, 2013). Thyroid hormone is known to play an important role in skull morphogenesis and multiple cranial malformations are associated with aberrant TH signaling (Desjardin et al., 2014; Hanken & Hall, 1988; Hanken & Summers, 1988; Hirano, Akita, & Fujii, 1995).

Thyroid hormone exerts extensive effects on the growth and remodeling of both endochondral and intramembranous bone by regulating gene expression (Bassett & Williams, 2003; Gogakos, Bassett, & Williams, 2010; Harvey et al., 2002; Waung, Bassett, & Williams, 2012; Wojcicka et al., 2013). The membranes of most skeletal cells possess TH transporters that facilitate hormone uptake (Gogakos et al., 2010). Active (T3) and inactive (T4) forms of TH are secreted by thyroid follicles and both are transported into bone cells where T4 is activated by cytoplasmic enzymes via the removal of an iodine molecule (Waung et al., 2012). Gene transcription is activated by T3 when it enters the nucleus and forms complexes with TH receptors bound to TH response elements in a number of target genes (Gogakos et al., 2010; Waung et al., 2012). The transcription products of these genes mediate the initiation and duration of bone matrix formation by regulating the differentiation and activity of bone-building cells (osteoblasts) and bone-resorbing cells (osteoclasts; Bassett & Williams, 2003; Harvey et al., 2002; Wojcicka et al., 2013).

In addition to the known roles of TH in both vertebrate metamorphosis and bone morphogenesis, changes to TH signaling may have been an important component of the diversification of the functional morphology of cypriniform feeding (McMenamin et al., 2017; Shkil & Smirnov, 2015; Shkil et al., 2015; Shkil, Kapitanova, Borisov, Abdissa, & Smirnov, 2012). To better understand the controls of fish metamorphosis and the developmental determinants of cypriniform jaw protrusion ability we measured the effects of different TH levels on the development of the functional morphology of feeding in the zebrafish. We also sought to determine whether modulating TH levels would alter zebrafish skull development in ways that could affect evolutionary potential. Variation in heritable traits facilitates evolutionary change and higher levels of variation tend to increase evolutionary potential (i.e., “evolvability”; Klingenberg, Duttke, Whelan, & Kim, 2012; Lavergne & Molofsky, 2007). Covariation among traits will facilitate their coordinated

evolution, but will tend to constrain independent evolutionary changes among subsets of covarying traits (Cooper, Wernle, Mann, & Albertson, 2011; Klingenberg, 2008; Sanger, Mahler, Abzhanov, & Losos, 2012; Young & Hallgrímsson, 2005). To gain insight into whether changes in TH levels could affect the evolutionary potential of fish skulls we compared both cranial shape variation and patterns of shape covariation between different skull regions among zebrafish that experienced a range of TH levels during development.

We collected morphological and functional data from a developmental range of zebrafish in which TH production was elevated (TH⁺), normal, or eliminated (TH⁻). We also collected morphological data from the premaxillary bones of nine additional species of danionin minnows (Danionini; Danioninae; Cyprinidae) that exhibit extensive diversity in adult jaw protrusion ability. Most cypriniform fishes possess a jaw protrusion mechanism that differs from those typically found in acanthomorphs in that they have a medial, mobile kinethmoid bone in the anterodorsal region of the skull that contributes motion to jaw protrusion (Gidmark, Staab, Brainerd, & Hernandez, 2012; Hernandez, Bird, & Staab, 2007; Motta, 1984). Although variation in jaw protrusion mechanisms have evolved among the ~20,000 cypriniform and acanthomorph fishes, in both clades the ascending arm of the premaxilla (sometimes referred to as the ascending process) typically slides anteriorly over the rostrum (the anterior portion of the neurocranium) during jaw protrusion and the length of the ascending arm is an important determinant of maximum protrusion distance (Cooper, Carter, et al., 2017; Motta, 1984).

We tested the following predictions: (a) normal TH levels are required for the development of functional abilities important to adult zebrafish feeding; (b) TH levels lead to alterations in the level of developmental variation in zebrafish head shape; (c) normal TH levels are required for the development of the wild-type pattern of covariation between different regions of the zebrafish skull; and (d) the functional morphology of jaw protrusion in adult TH⁻ zebrafish closely resembles that in related minnows with limited protrusion abilities.

2 | MATERIALS AND METHODS

2.1 | Study system

We utilized three zebrafish lines to study the effects of TH on the development of their feeding biomechanics: (a) the transgenic line *Tg(tg:nVenus-2a-nfnB)^{wp.rts}* in which the thyroid follicles can be chemically ablated; (b) the mutant line *opallus^{b1071}*, hereafter TH⁺, which has a missense mutation in thyroid stimulating hormone receptor (*tshr*) that causes constitutive hyperthyroidism (i.e., elevated TH levels; McMenamin et al., 2014); and (c) the AB wild-type line (euthyroid, i.e., normal TH). Both the transgenic and the mutant lines originated in the AB line (McMenamin et al., 2014). All fish were maintained under standard conditions at 28°C and a 14-hr light/10-hr dark cycle under approved WSU IACUC Protocol 04285.

Hypothyroid specimens (TH⁻) were produced via nitroreductase-mediated cell ablation of thyroid follicles in *Tg(tg:nVenus-2a-nfnB)^{wp.rts}* larvae at 4 days postfertilization (dpf) following McMenamin et al. (2014). Ablation was performed immediately after formation of the thyroid follicles so that they were rendered permanently incapable of hormone

production. In brief, TH-fish were produced by placing specimens in a solution of 1% dimethyl sulfoxide (DMSO) and 10 mM metronidazole for 4 hr. Control *Tg (tg:nVenus-2a-nfnB)^{wp.rt8}* fish treated with 1% DMSO only for 4 hr will be referred to as “DMSO” hereafter.

All specimens were fed live paramecia exclusively from 5 dpf until they were large enough to begin feeding on live *Artemia*. *Artemia* were introduced slowly into the diet from ~14dpf until all specimens in an aquarium were capable of consuming brine shrimp (as indicated by exhibiting pink, distended bellies after feeding), at which point paramecia were no longer used. To provide more nutritionally complete *Artemia*, newly hatched brine shrimp were collected after 24 hr and fed for an additional 24 hr with an infusion of *Spirulina* sp. algae (RGcomplete, Reed Mariculture, Inc., Campbell, CA). To prevent exposure to exogenous TH present in most prepared fish foods, all fish were fed exclusively with enriched *Artemia* after they were large enough to be fully weaned off of paramecia. All specimens were maintained in 9 L aquaria from 5 dpf onward on a recirculating system with carbon filters. Specimens from each treatment were sampled for kinematic and shape analyses at 8, 15, 30, 60, and 100 dpf.

2.2 | Kinematic analyses

To test our first prediction that normal TH levels are required for the development of functional abilities important to adult zebrafish feeding we used high speed video to measure multiple aspects of feeding performance in zebrafish from all TH treatment groups. Fish were filmed in lateral view while feeding on either paramecia (8 and 15 dpf) or *Artemia* (30, 65, and 100 dpf). We analyzed kinematic data from five feeding strikes for each individual and examined five specimens of each age class from each treatment group. Feeding strikes were recorded at 500 frames/second using an Edgertronic monochrome high-speed video camera (Sanstreak Corp., San Jose, CA). Kinematic analyses of feeding strikes were performed using the ImageJ software program (Schneider et al., 2012). We measured the following variables in each video frame of all feeding strikes (see Figure 1a for reference): gape angle (the angle created between landmarks A, B, and C with B as the vertex), jaw protrusion (linear distance between landmarks A and F minus the minimum distance recorded between A and F for that strike), hyoid depression (linear distance between landmarks D and E minus the minimum distance recorded between D and E for that strike), and cranial elevation angle (the angle created by the intersection of a line running along the dorsal edge of the head with a line running along the dorsal edge of the body, with landmark F denoting the point of head rotation). All measurements were made by the same researcher to minimize introduction of operator error. Jaw protrusion and hyoid depression were standardized by fish standard length (SL). Maximum values for every variable were recorded for each feeding strike of every specimen. These maxima were used to calculate a mean value for each specimen. For each age class we used analysis of variance (ANOVA) to test for differences between treatment groups for the maximum value of each variable. When a significant difference was detected we used a Tukey’s honestly significant difference (HSD) test to determine which treatment groups were significantly different from each other.

2.3 | Morphological analyses

The specimens used in feeding trials were also used for morphological analyses. Euthanized specimens were fixed in 4% formalin for 24 hr and then stepped over into 70% ethanol. They were then cleared and stained for bone and cartilage. The smaller specimens (8–30 dpf) were processed according to Walker and Kimmel's (2007) acid-free staining protocol. Larger specimens (65 and 100 dpf) were cleared and stained according to Potthoff (1984). Specimens were then stepped into 80% glycerol and photographed in lateral view using an Olympus DP25 digital camera interfaced with an Olympus SZ61 dissecting microscope.

The program tpsDIG2 (<http://life.bio.sunysb.edu/morph/>) was used to place landmarks (LM) and semi-landmarks (semi-LM) on digital images of fish heads (Figure 1b,c). We chose skeletal LM that are present at all of the developmental stages examined. Semi-LM were used to capture the shape of curved surfaces LM (Figure 1b,c). The programs tpsUtil and tpsrelW (<http://life.bio.sunysb.edu/morph/>) were then used to superimpose semi-LM using a chord-distance (Procrustes distance) based “sliders” method and to remove size and orientation differences from LM and semi-LM position data via Procrustes transformations. Pooled shape data from all specimens of all ages in each hormonal treatment group ($N = 25$ for each treatment) were used to: (a) test for differences in head shape variation over the course of development; and (b) test for differences in the patterns of covariation between LM and semi-LM locations over the course of development.

To test our second prediction that changes in TH levels will affect developmental variation we measured head shape variation in each of our TH treatment groups. We then used a resampling approach to test for differences in shape variation. Head shape variation was quantified by calculating the Foote disparity value for each treatment group (Foote, 1993). A permutation procedure (2,000 iterations) was used to test for differences in disparity between pairs of data sets. If actual differences in shape disparity values were greater than the upper bound of a 95% confidence interval calculated via permutation, then the disparity values were considered to be significantly different. We used DisparityBox, which is an analytical tool available within the PCAGen8 program, to perform these calculations.

We used a principal components based resampling method to test our third prediction that normal TH levels are required for the development of the wild-type pattern of covariation between different regions of the zebrafish skull. We performed principal components analysis (PCA) of LM and semi-LM positions using the program PCAGen8. These PCAs utilized covariation matrices that capture patterns of positional covariation. To determine if patterns of covariation were significantly different between the hormonal treatments we conducted pairwise comparisons of PC axis orientations using a bootstrapping procedure (4,900 sets).

Although the different PC axes derived from the same data are orthogonal to each other, they are not independent, since only those aspects of covariation that were not associated with PC1 can be used to define subsequent axes. It therefore impossible to compare the orientations of PC axes subsequent to PC1 individually. All analyses that involved multiple axes determined whether the alignments of planes (2 axes) or multidimensional hyperplanes (3 axes) were significantly different.

The shape of the premaxillary bone in the upper jaw is a major determinant of maximum jaw protrusion distance in fishes (Cooper, Carter, et al., 2017; Motta, 1984). We used a principal components based resampling method to test our fourth prediction that the functional morphology of jaw protrusion in adult TH⁻ zebrafish closely resembles that in related minnows with limited protrusion abilities. We compared developmental variation in the premaxillary shape of AB and TH⁻ zebrafish to the variation in adult premaxillary shape that has evolved among other members of the cyprinid tribe Danionini (sensu Tang et al., 2010). We obtained specimens of the following fishes through the pet trade: *Danio albolineatus*, *Danio erythromicron*, *Danio feegradei*, *Danio kyathit*, *Danio nigrofasciatus*, *Danionella translucida*, *Devatio aequipinnatus*, *Devatio maetaengensis*, and *Mi-crodevatio kubotai*. Adult specimens (2–4 per species) were cleared and stained following Potthoff (1984). AB zebrafish at 35, 65, and 100 dpf were also cleared and stained (five specimens per age class). We used 35 dpf specimens instead of 30 dpf fish because some 30 dpf specimens did not have well-ossified premaxillae. Premaxillae were removed from all specimens after clearing and staining and then photographed as described above. Anatomical LM and semi-LM (Figure 4b,c) were used to quantify premaxillary shape. PC scores were used to calculate a developmental trajectory for premaxillary shape change in AB zebrafish.

For all shape analyses we used the program Coord-Gen8 to transform the LM and semi-LM coordinate data from the format used by the tpsDIG2 program into the format utilized by Mac OS (Apple Inc.) versions of the IMP-8 series of programs. The IMP-8 programs Coord-Gen8, PCAGen8, TwoGroup8, and PCAGen8 were developed by David Sheets and are available for download at: <http://www3.canisius.edu/~sheets/IMP8.htm>.

3 | RESULTS

3.1 | Ossification patterns

All specimens possessed largely cartilaginous crania at 8 and 15 dpf. Skulls were mostly, but not completely, ossified by 30 dpf in specimens from all treatment groups except TH⁻ (Figure 2). We will refer to 8 and 15 dpf specimens as “premetamorphic,” 30 dpf specimens as “mid-metamorphic” and 65 and 100 dpf specimens as “postmetamorphic.” Euthyroid and TH⁺ fish had fully ossified skulls by 65 dpf, while TH⁻ specimens retained cartilaginous regions in the calvarium (skull vault) at this stage and in some TH⁻ fish this region failed to ossify by 100 dpf (Figure 2).

3.2 | Standard length

Fish SLs did not differ significantly between the AB, DMSO, and TH⁺ treatments at any developmental stage (Figure 3). Note that since age alone is a limited proxy for development (Parichy, 2009; McMenamin, Chandless, & Parichy, 2016), individuals within these age categories likely represented a range of developmental stages. Nonetheless, SL did not differ significantly different for any treatment before 65 dpf. At 65 and 100 dpf, TH⁻ fish were shorter than those in other treatments, suggesting TH is required for normal growth rates (Figure 3).

3.3 | Jaw morphology

The lower jaws of TH⁺ specimens began to show abnormal enlargement by 30 dpf (Figures 2 and 3). Lower jaw morphology was normal in TH⁻ specimens (Figures 2 and 3). The maxillary and premaxillary bones of the upper jaw were smaller in 65 and 100 dpf TH⁻ zebrafish relative to both euthyroid and TH⁺ specimens (Figure 2). Upper jaw morphology in TH⁺ specimens appeared comparable to that in euthyroid specimens (Figure 2).

3.4 | Kinematic analyses

Three kinematic variables could be measured at all of the developmental stages examined (cranial elevation, gape distance and hyoid depression). At each developmental stage these three variables were compared among all treatment groups (six comparisons per variable). In regard to these three variables there were therefore 36 comparisons of kinematic performance both before (8 and 15 dpf) and after (65 and 100 dpf) metamorphosis. Of the 36 premetamorphic comparisons, 11 of them exhibited significant differences (30.6%), while there were significant differences between 13 of the 36 postmeta-morphic comparisons (36.1%; Table 1). Treatment groups were most similar in regard to cranial elevation (five significant differences total) and most different in regard to gape angle (13 significant differences total; Table 1). Fish from the euthyroid treatments (AB and DMSO) showed the most similarity in cranial movement, with only three significant kinematic differences between these two treatments (all developmental stages combined; Table 1). Fish from the TH⁻ treatment exhibited the most limited range of cranial element motion during feeding (Table 1 and Figures 5 and S2). There were also 22 significant kinematic differences (all developmental stages combined) between TH⁻ specimens and all other treatment groups, which was the highest number for any of the treatment groups (Table 1).

Maximum gape angle occurred at or near time 0 for all treatment groups throughout development (Figure 5a,b and S2). AB fish exhibited high gape angles throughout development (Figures 5a,b and S2), while TH⁻ fish exhibited low gape angles throughout development (Figures 5a,b and S2). Maximum hyoid depression occurred after time 0 for all treatment groups throughout development (Figure 5c,d and S2). Hypothyroid fish exhibited low levels of hyoid depression throughout development until 100 dpf, when they displayed a high degree of hyoid depression (Figures 5c,d and S2). Maximum cranial elevation occurred at or immediately after time 0 throughout development (Figures 5e,f and S2). Hypothyroid fish exhibited low levels of cranial elevation throughout development (Figures 5e,f and S2). Measureable upper jaw protrusion was not observed before 30 dpf in any treatment group, but a small number of euthyroid and TH⁺ specimens exhibited marginal upper jaw movement at this stage. Specimens from all treatment groups except TH⁻ exhibited protrusile upper jaws by 65 dpf. In contrast, hyperthyroid fish demonstrated low levels of jaw protrusion at both 65 and 100 dpf and were significantly different from all other treatment groups at both stages (Table 1 and Figure 5g,h).

3.5 | Developmental variation in head shape

Hypothyroid fish showed significantly less developmental variation in head shape than any other treatment group (Table 2). Hyperthyroid fish showed significantly greater developmental variation in head shape than both TH⁻ and DMSO specimens.

3.6 | Shape covariation

Patterns of shape covariation were highly similar in the skulls of all treatments with one exception: TH⁻ versus TH⁺. These two treatments produced zebrafish skulls that exhibited similar patterns of shape covariation in that the orientations of their PC1 axes were not found to be significantly different from parallel to one another (Table 3). However, their PC2 axes were not parallel, which indicates a moderate difference in anatomical trait covariation (Table 3). Less pronounced differences in TH levels did not produce detectable differences in patterns of skull shape covariation. The first four PC axes described similar covariation patterns when the skulls of euthyroid fish (both AB and DMSO) were compared to those of either TH⁻ or TH⁺ specimens (Table 3). The first four PC axes also described similar covariation patterns in skulls from the two euthyroid treatments (Table 3).

3.7 | Head and premaxilla shape

Head shape differences between TH treatments were present at all ages examined, but those from fish in different treatments were most similar at the youngest ages (8 and 15 dpf; Figure S1). By 100 dpf TH⁺ and TH⁻ head shapes were highly distinct from each other (Figure S1).

The ascending arm of the premaxilla (Figure 1), is extremely short at 35 dpf in AB fish, but elongates significantly by 100 dpf (Figure 4). Premaxilla shape in 100 dpf TH⁻ specimens was similar to that of newly ossified premaxillae in 35 dpf AB fish, in that they had very short ascending arms (Figure 4). Both the maxillae and premaxillae in the upper jaws of TH⁻ fish exhibited limited growth in size that was unaccompanied by the shape changes seen in those of postmetamorphic euthyroid and TH⁺ fish (Figures 2 and 4).

Variation in ascending arm length was strongly associated with both postmetamorphic AB zebrafish development and danionin evolution (Figure 4). Developmental variation in AB premaxilla shape was not significantly different from the variation in premaxilla shape that has evolved among nine additional danionin species (i.e., the first 4 PC axes derived from these two data sets were parallel; Table 3). Among the species that we examined, those from the genus *Danio* have longer premaxillary ascending arms than those from other danionin genera, with *Danio erythromicron* having the longest arms (Figure 4). The premaxillae of *Danionella*, *Devatio*, and *Microdevatio* were most similar to those from 35 dpf AB and 100 dpf TH⁻ zebrafish. Manipulation of freshly euthanized specimens from these three genera, as well as observations of their feeding using high-speed video (personal observation; S. McMenamin et al., 2017), indicate that jaw protrusion is extremely limited to nonexistent in these species.

4 | DISCUSSION

We found strong evidence that TH levels affect the development of zebrafish feeding biomechanics (prediction 1), that TH increases developmental variation in zebrafish head shape (prediction 2), and that the functional morphology of jaw protrusion in adult TH⁻ zebrafish duplicates that of closely related minnows with limited protrusion abilities (prediction 4). Normal TH levels do not appear to be required for the development of the

wild-type pattern of covariation between different regions of the zebrafish skull (prediction 3), but specimens with highly divergent TH levels (TH⁻ and TH⁺) did exhibit limited differences in cranial covariation patterns. We also found evidence that the developmental trajectory of premaxillary shape change in AB zebrafish is parallel to an important evolutionary axis of danionin diversification (premaxilla shape and jaw protrusion ability).

We found that differences in feeding biomechanics between our TH treatment groups were present at all of the developmental stages we examined, but that these differences increased with age and were more pronounced after metamorphosis (Table 1 and Figure S1). The presence of TH appears to be particularly important for the transition from larval to adult feeding mechanics in the zebrafish. An absence of TH delayed cranial ossification and arrested premaxillae formation so that upper jaw protrusion was absent or severely limited in postmetamorphic TH⁻ zebrafish development to permit upper jaw protrusion (Table 1 and Figures 2, 4, and 5; S. McMenamin et al., 2017).

Hypothyroidism appeared to truncate aspects of zebrafish cranial development. In addition to arresting premaxilla morphogenesis immediately after ossification (Figure 4), lack of TH caused a significant reduction in the skull shape variation that arose between 8 and 100 dpf (Table 2). However, TH⁻ fish grow more slowly than the other groups (Figure 3) and some of this decrease in variation may be attributable to TH⁻ specimens' slow growth or narrow range of developmental stages. Conversely, TH⁺ fish showed an increase in developmental variation in skull shape relative to both DMSO and TH⁻, though they were not significantly different from AB specimens in this regard (Table 2).

The only significant differences in cranial shape covariation that we observed were between the TH⁻ and TH⁺ treatments, but these differences did not include PC1, which is the largest axis of shape variation (Table 3). Changes to patterns of anatomical trait covariation can have debilitating effects, especially in biomechanical systems where there is a high level of functional integration between different elements (Armbruster, Pelabon, Bolstad, & Hansen, 2014; Kimmel et al., 2015; Walker, 2007) and both TH⁻ and TH⁺ fish retained a sufficient level of integration between the various bones of the skull to feed successfully.

Premaxillary ascending arm length is an important determinant of maximum upper jaw protrusion distance (Figure 1; Cooper, Carter, et al., 2017; Hulsey, Hollingsworth, & Holzman, 2010; Motta, 1984). Eliminating TH produces zebrafish with highly reduced ascending arms and feeding mechanisms that resemble those of close relatives with nonprotrusile jaws (Table 1 and Figures 4 and 5; S. McMenamin et al., 2017). Many of these have been described as being paedomorphic (Britz, Conway, & Rueber, 2014; Kullander & Britz, 2015), a condition that is strongly associated with changes in TH signaling in vertebrates (De Groef, Grommen, & Darras, 2018; Ivanovic, Cvijanovic, Denoel, Slijepcevic, & Kalezic, 2014; Laudet, 2011). Differences in jaw protrusion ability are strongly associated with differences in diet in marine damselfishes (Cooper, McGraw, Khazanchi, & Ieee., 2017), and although diet data for wild danionins is limited or nonexistent, we suspect that the same may be true for the species examined here. Although TH⁻ zebrafish did not develop adult jaw protrusion abilities, they retained sufficient functional integration to be able to feed successfully in the lab environment. It has been

suggested previously that evolutionary modulation of TH signaling may have contributed to the diversification of cypriniform fishes (McMenamin et al., 2017; Shkil & Smirnov, 2015; Shkil et al., 2012, 2015). We conclude that evolutionary changes in TH signaling may be capable of inducing some adaptive changes in the development of fish feeding mechanics without causing severe disruptions to cranial morphogenesis that produce nonfunctional skulls.

4.1 | The importance of late developmental periods to fish evo-devo

The field of evo-devo is focused on understanding the connections between developmental processes and evolutionary change (Carroll, 2008). Phylogenetic analyses of comparative data have expanded tremendously in recent years and the evolutionary patterns traced by many lineages have been described in great detail (Freckleton, Harvey, & Pagel, 2002; Garland, Bennett, & Rezende, 2005; Mouquet et al., 2012). A particularly rich source of comparative information exists for the field of fish feeding biomechanics and we now know a great deal about the ecological, morphological, and functional evolution of many fish clades. Due to the use of multiple fish species as model organisms for developmental study (e.g., zebrafish, medaka, Mexican tetra, fugu, stickleback, multiple African rift-lake cichlids, etc.) it is possible to experimentally explore aspects of fish development that have played important roles in shaping evolutionary diversification. However, since most evolutionary studies of fish feeding have focused on adult specimens and since most fishes do not develop adult feeding biomechanics until late in their development, often after a pronounced metamorphosis, merging these two areas of investigation will require an increased focus on the later stages of skull morphogenesis.

Evo-devo studies of jaw protrusion provide an illustration of this point. Protrusile jaws constitute one of the most significant biomechanical innovations to arise in fish skulls (Motta, 1984; Wainwright & Longo, 2017; Wainwright et al., 2015), but we know of no species in which protrusion arises before the larva-to-juvenile transition. The fact that small fish larvae will experience much higher water viscosities than larger adult forms (Hernandez, 1995; Hernandez, 2000) may predispose many species to undergo developmental changes in feeding mechanics. We suggest that this may be particularly true for fishes that utilize jaw protrusion during adult feeding. Thyroid hormone signaling has pervasive effects on skeletal morphogenesis during late development (Okada, Tanaka, & Tagawa, 2003; Pascual & Aranda, 2013; Shkil et al., 2012), spikes in TH are associated with metamorphosis of larval fishes into juveniles (McMenamin & Parichy, 2013; Okada et al., 2003), and adult jaw protrusion abilities do not arise in TH⁻ zebrafish (Table 1, Figure 4, and 5; S. McMenamin et al., 2017). The range of effects that TH has on bone remodeling (Bassett & Williams, 2003; Gogakos et al., 2010; Harvey et al., 2002; Waung et al., 2012; Wojcicka et al., 2013) suggests that there are likely to be a number of mechanisms by which changes to TH signaling could alter the development of fish skulls and jaws. Our data support the assertion that a better understanding of how TH affects cranial remodeling has important relevance to the evo-devo of adaptive diversification in fish feeding.

Evolutionary changes to TH signaling may have contributed to the diversification of cyprinid fishes. Miniaturization has arisen multiple times in danionine minnows (e.g., the genera

Danionella, *Fangfangia*, *Microdevario*, *Paedocypris*, and *Sundadanio*) and it has been speculated that developmental truncation (i.e., paedomorphosis) has contributed to the divergence of these species from their ancestral forms (Britz, Conway, & Ruber, 2009; Britz et al., 2014; Britz, Kottelat, & Tan, 2011; Ruber, Kottelat, Tan, Ng, & Britz, 2007). We observed that loss of TH production retarded zebrafish growth (Figure 3) and arrested premaxillary shape development at an early stage so that adult TH⁻ specimens did not develop adult jaw protrusion abilities (Table 1, Figures 4 and 5). McMenamin et al. (2017) demonstrated that adult TH⁻ zebrafish retained feeding mechanics similar to those of both larval euthyroid zebrafish and adult specimens from the highly paedomorphic danionine genus *Danionella* (Britz et al., 2009). Shkil et al. have provided evidence consistent with the hypothesis that changes in TH production and/or tissue responsiveness to TH has contributed to heterochronic evolution during the adaptive divergence of cyprinid fishes outside of the danionine lineage (Shkil & Smirnov, 2015; Shkil et al., 2012, 2015). Taken together, these findings suggest that evolutionary changes in TH signaling may have shaped aspects of cypriniform diversification, especially during the repeated evolution of miniature species that exhibit heterochronic development in which they retain aspects of larval feeding mechanics as adults.

5 | CONCLUSIONS

TH signaling induced development of an adult premaxillary shape and adult feeding kinematics in metamorphosing zebrafish. Hypothyroidism inhibited the development of jaw protrusion by severely reducing the development of premaxillary ascending arm length. Normal ontogenetic changes in zebrafish premaxillary shape represent developmental variation that could underlie evolutionary changes in jaw protrusion ability. The pronounced effects of thyroid hormone on the development of the functional morphology of the skull during the juvenile to adult transition suggest that changes in TH or pathways affected by TH may have contributed to adaptive diversification of fish feeding biomechanics.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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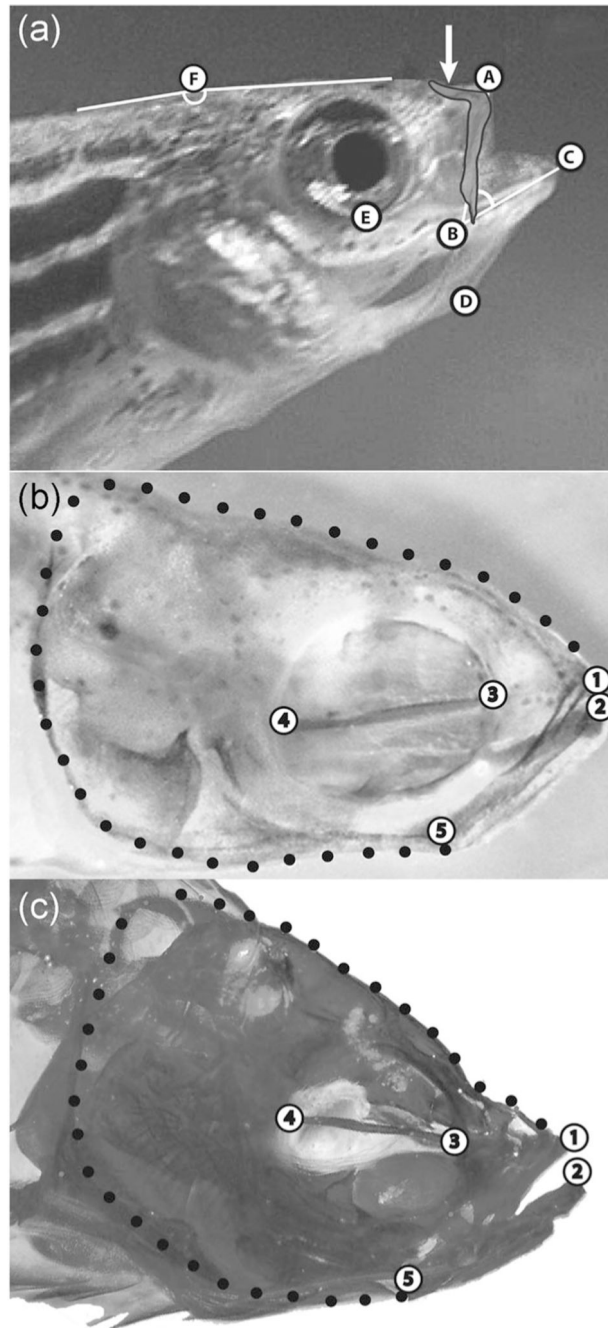


FIGURE 1.

Landmarks and semilandmarks used in analyses of movement and shape. (a) Landmarks used in kinematic analyses with an image of the premaxillary bone of the upper jaw superimposed its correct anatomical position in a fully protruded upper jaw. Arrow indicates the ascending arm of the premaxilla. Landmarks: (A) Anterior tip of the upper jaw; (B) corner of the mouth; (C) anterior tip of the lower jaw; (D) anterior tip of the hyoid; (E) ventral-most point of the orbit; and (F) vertex of the angle used to measure head rotation during cranial elevation (the dorsal surface of the head anterior to this point rotated upward

during cranial elevation when feeding, while the dorsal surface of the trunk posterior to this point did not). (b) Anatomical landmarks and semilandmarks used in shape analyses of all specimens at all ages sampled shown on a larval zebrafish. Landmarks: (1) anterior tip of the premaxilla in the upper jaw; (2) anterior tip of the dentary bone in the lower jaw; (3) junction of the parasphenoid with the anterior wall of the orbit; (4) junction of the parasphenoid with the posterior wall of the orbit; and (5) lower jaw joint (articularquadrate joint). Black circles indicate semi-landmarks evenly spaced between LM 1 and 2 to capture overall head shape. (c) Anatomical landmarks and semi-landmarks used in shape analyses of all specimens at all ages sampled shown on an adult zebrafish (LM are the same as those in panel b). LM, landmark

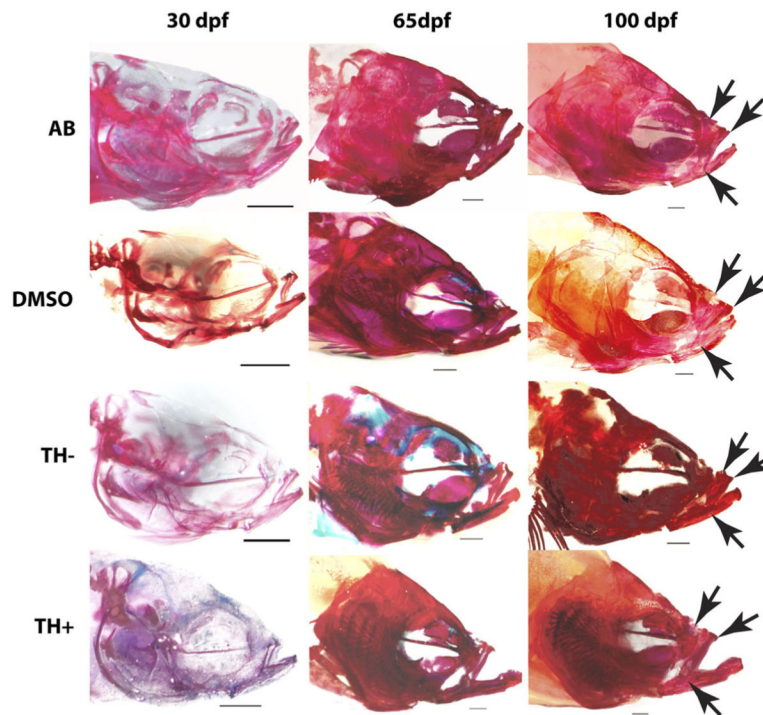
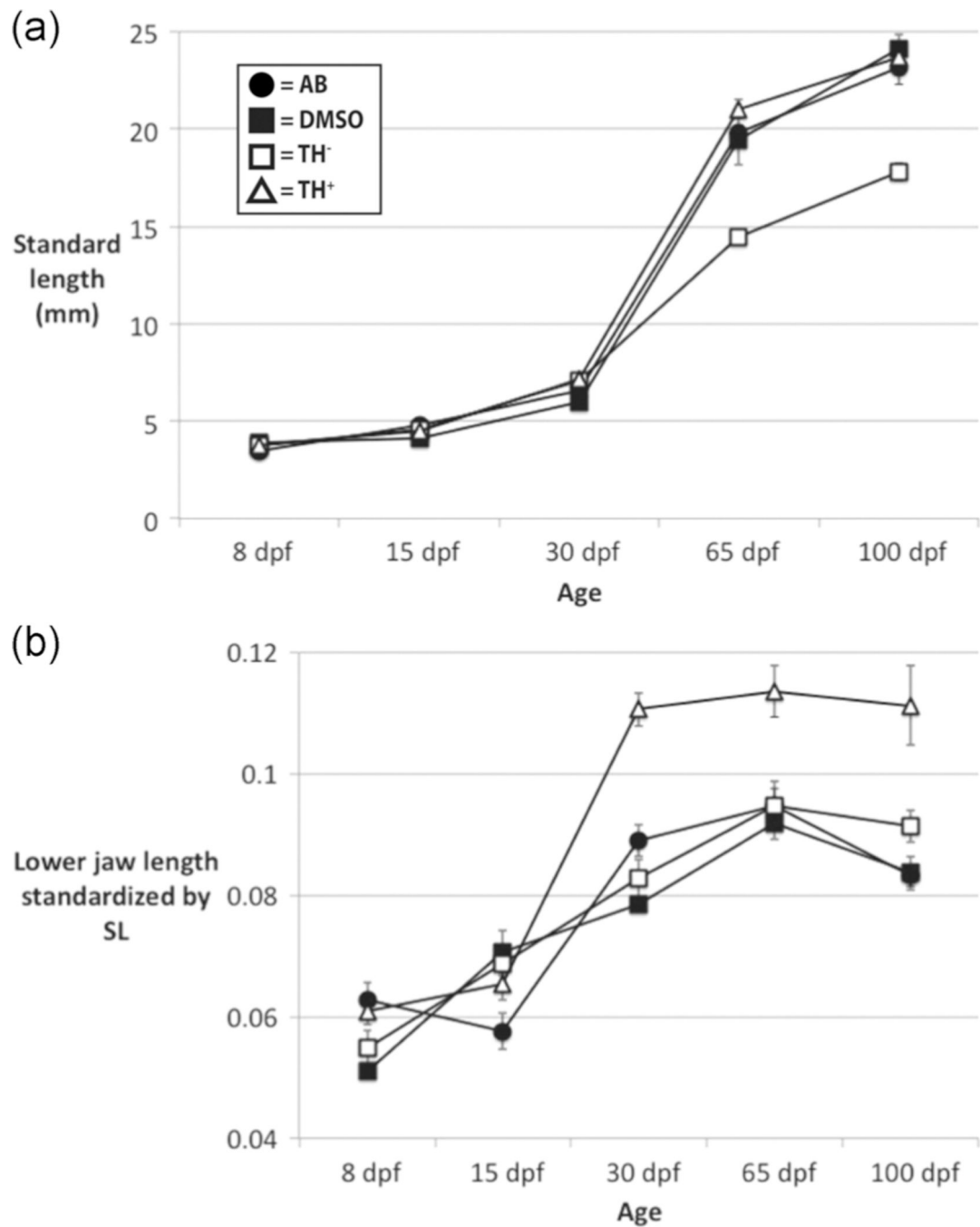
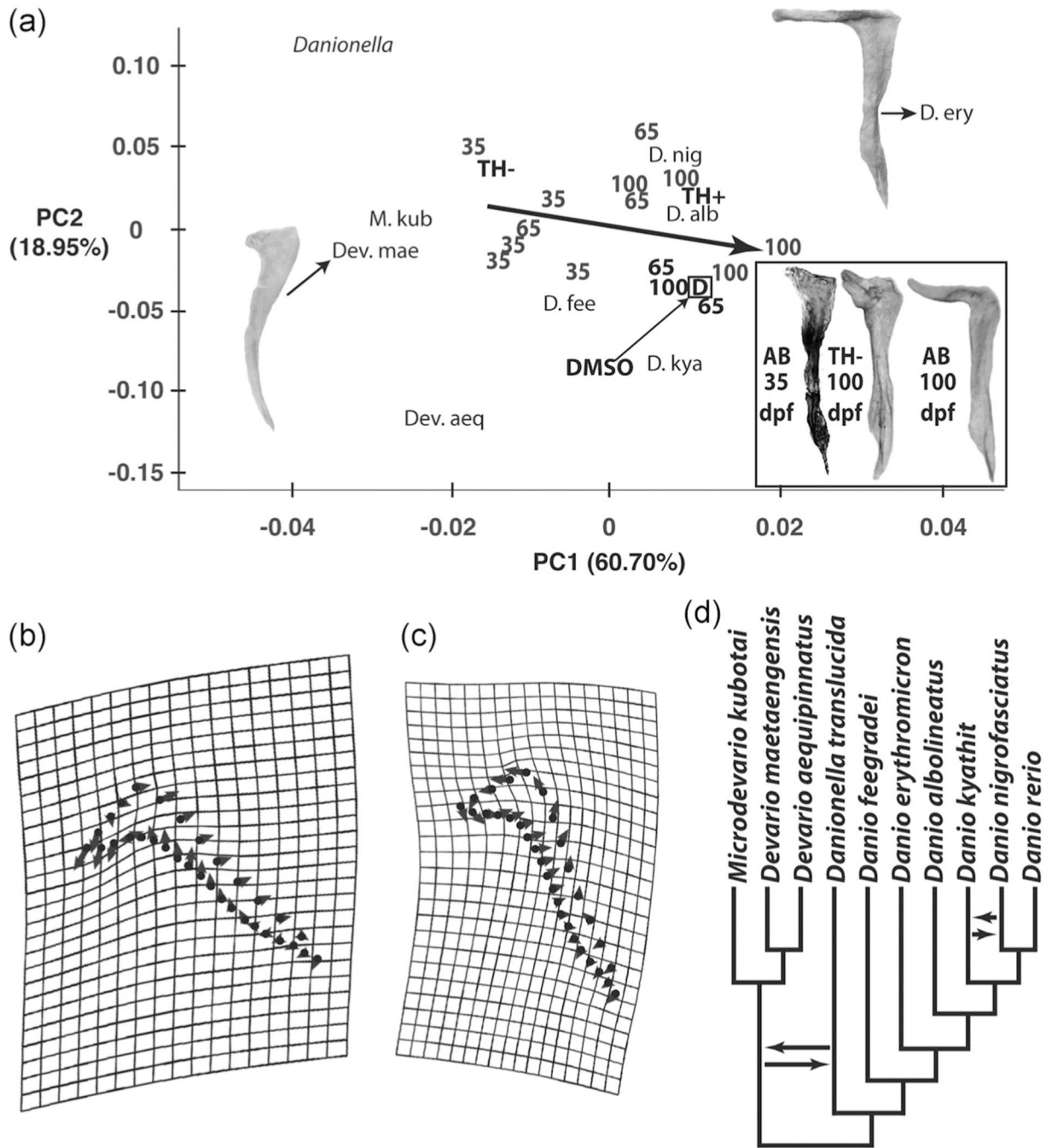


FIGURE 2.

Representative cleared and stained specimens of postmetamorphic specimens from all treatment groups. Blue coloration indicates cartilage stained by alcian blue. Red indicates bone stained by alizarin red. Arrows represent the antero-dorsal and postero-dorsal edges of the ascending arm of the premaxillary bone. Arrowheads indicate the postero-ventral tip of the dentigerous arm of the premaxilla (scale bars = 1 mm). TH⁻ specimens exhibited delayed cranial ossification. In many of our TH⁻ specimens the calvarium (skull vault) was not fully ossified by 100 dpf. Both the overall size of the premaxilla and the length of its ascending arm relative to the length of its dentigerous arm were reduced in TH⁻ fish. TH⁺ fish exhibited hypertrophied lower jaws. TH, thyroid hormone

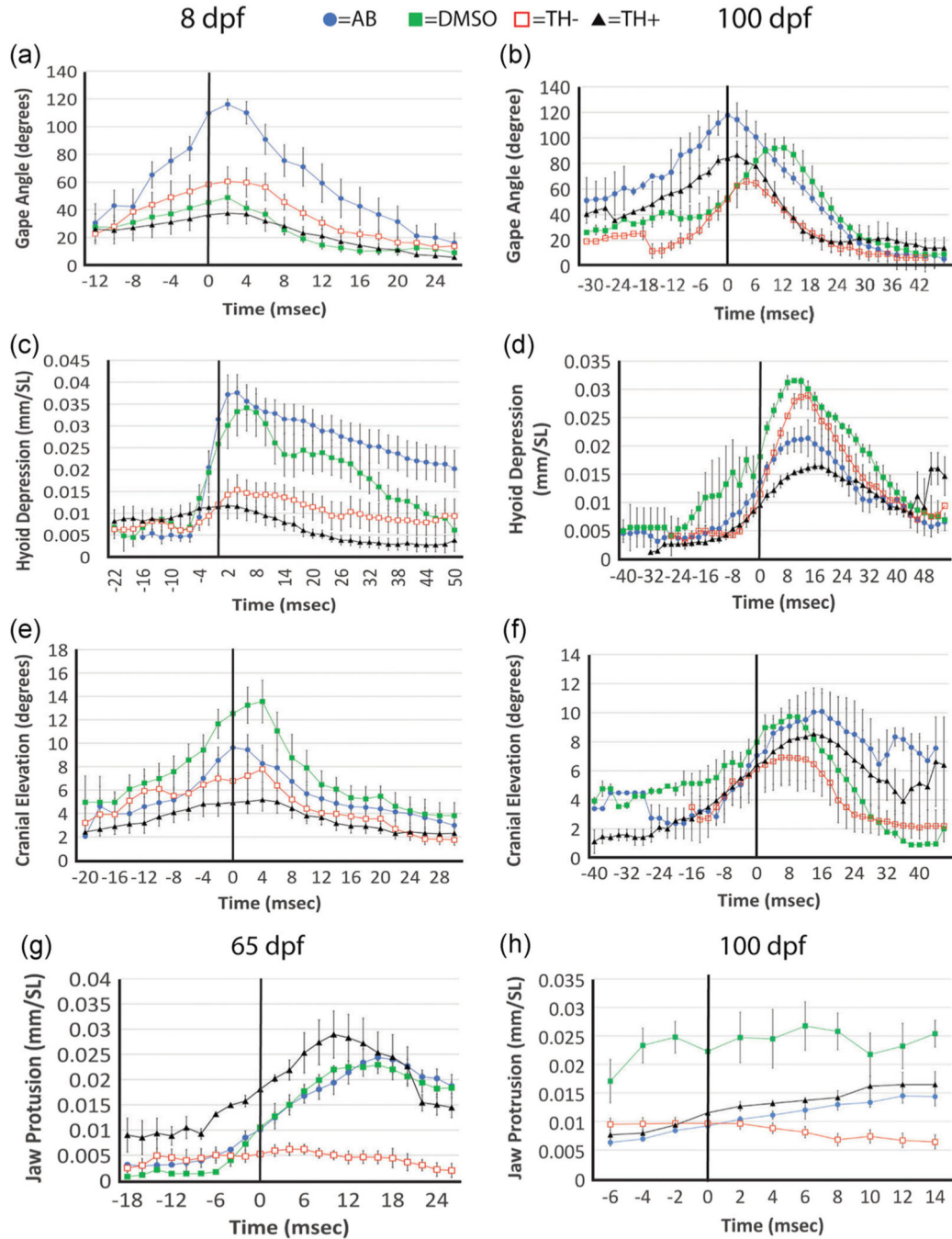
**FIGURE 3.**

Comparisons of anatomical growth among treatment groups. Mean sizes with standard error bars are given for all treatments at each age sampled. Key to symbols in panel (a). (a) All treatment groups exhibited similar increases in body length until 30 dpf, after which TH⁻ fish exhibited lower growth rates. (b) All treatment groups exhibited similar increases in the relative length of the lower jaw until 15 dpf. After 15 dpf the lower jaws of TH⁺ fish grew much more quickly than those of the other treatment groups and then stabilized at a larger relative size. TH, thyroid hormone

**FIGURE 4.**

Comparisons of premaxillary morphology. (a) Principal component score plot derived from coordinate-based analyses of premaxilla shape in the following specimens: (1) 35, 65, and 100 dpf AB zebrafish; (2) 100 dpf DMSO, TH⁻, and TH⁺ zebrafish; and (3) adults of nine additional species of danionin minnows. The difference in premaxillary shape between *Devario maetaeangensis* and *Danio erythromicron* exemplify the shape diversity explained by PCI. PC1 is strongly associated with the length of the ascending arm relative to the length of the dentigerous arm. Numbers indicate the distributions of 5 specimens of AB zebrafish

from each of the three ages sampled (numbers correspond to dpf). The location of the Procrustes mean premaxillary shapes of TH⁺ and TH⁻ specimens are indicated by their respective symbols. The location of the Procrustes mean premaxillary shape of DMSO specimens is indicated by “D”. The location of Procrustes mean shapes of the nine nonzebrafish premaxillae are indicated by an abbreviation of the scientific name of each species (see panel (d) for full species names), except for *Danionella translucida*, where the complete genus name is used. Images of premaxilla shapes of particular interest are inserted. The largest arrow denotes the developmental shape trajectory for AB premaxillae. (b) Landmarks used in shape analyses (black circles), deformation grid and vector plot that shows the shape variation associated with PC1. (c) Landmarks used in shape analyses (black circles), deformation grid and vector plot that shows the shape variation associated with PC2. (d) Phylogenetic relationships of the 10 species whose premaxillae were compared. The phylogeny depicted is taken from Tang et al. (2010). Arrows indicate branch positions that are swapped in the relationships reported by McCluskey and Postlethwait (2015), who did not examine *Devario maetaeensis*. DMSO, dimethyl sulfoxide; PC, principal component; TH, thyroid hormone

**FIGURE 5.**

Comparisons of cranial movements during feeding among the four TH treatment groups. In all cases time zero represents the time point when live prey (paramecia at 8 dpf and brine shrimp nauplii at 65 and 100 dpf) passed the tips of the upper and lower jaws as they were being engulfed. Units and standardizations are given in parentheses in each case. (a-h) Plots for gape angle (a,b), hyoid depression (c,d) and cranial elevation (e,f) for specimen of ages 8 and 100 dpf, respectively (kinematic plots for these four variables at 15, 30, and 65 dpf are

presented in Figure S2). (g) Jaw protrusion at 65 dpf. (h) Jaw protrusion at 100 dpf. SL, standard length; TH, thyroid hormone

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ANOVA and Tukey's HSD results for comparisons of kinematic variables among treatment groups for each developmental stage sampled

TABLE 1

	<u>Premetamorphic</u>		<u>Midmetamorphic</u>		<u>Postmetamorphic</u>	
	8 dpf	15 dpf	30 dpf	65 dpf	100 dpf	
Gape angle	AB > DMSO	AB > TH ⁻	TH ⁺ > DMSO	AB > TH⁻	AB > DMSO	
	AB > TH⁻	AB > TH⁺	TH ⁺ > TH ⁻	DMSO > TH ⁻	AB > TH⁻	
	AB > TH⁺				DMSO > TH ⁻ TH ⁺ > AB	
Hyoid depression	AB > TH ⁻	AB > TH ⁻	AB > DMSO		DMSO > AB	
	AB > TH ⁺		AB > TH ⁻	ANOVA	DMSO > TH⁺	
	DMSO > TH ⁻		TH⁺ > DMSO	not sig.	TH ⁻ > TH ⁺	
Cranial elevation	DMSO > TH ⁺		TH ⁺ > TH ⁻			
	DMSO > TH ⁺		TH ⁺ > TH ⁻	AB > DMSO		
		ANOVA		AB > TH⁻	ANOVA	
Jaw protrusion		not sig.		TH⁺ > DMSO	not sig.	
				TH ⁺ > TH ⁻		
	NA	NA	NA	AB > TH⁻	AB > TH⁻	
			DMSO > TH ⁻	DMSO > TH⁻		
			TH⁺ > TH⁻	TH⁺ > TH⁻		

Note: Significant Tukey's HSD post-hoc comparisons are listed if the ANOVA results were significant ($\alpha = .05$). Bold/shaded comparisons had $p < .001$. The treatment with the higher maximum value is indicated for each comparison.

Abbreviations: ANOVA, analysis of variance; DMSO, dimethyl sulfoxide; dpf, days postfertilization; HSD, honestly significant difference; TH, thyroid hormone.

TABLE 2

Shape disparity comparisons for head morphology

	AB	DMSO	TH ⁻
ABFD = 0.00622046	NA		
DMSOFD = 0.00595240	DD = 0.000288	NA	
	UBCI = 0.00132652		
TH ⁻ FD = 0.00410838	DD = 0.002112 (AB)	DD = 0.001824 (DMSO)	
	UBCI = 0.00139538	UBCI = 0.00134556	NA
TH ⁺ FD = 0.00780772	DD = 0.001587	DD = 0.001875 (TH⁺)	DD = 0.003699 (TH⁺)
	UBCI = 0.00182215	UBCI = 0.00174844	UBCI = 0.00188894

Abbreviations: DD, difference in the disparities of the treatments compared; FD, Foote disparity; UBCI, upper bound of 95% CI. Disparities were significant when DD > UBCI. Treatments with higher disparities are indicated when significant (bold/shaded).

TABLE 3

Shape space orientation comparisons for head and premaxilla morphology

	Angle (degrees)	95% CI
PC1		
AB vs. DMSO	21.78	16.35–31.81
AB vs. TH ⁻	34.73	29.24–44.89
AB vs. TH ⁺	23.71	19.65–31.17
DMSO vs. TH ⁻	18.26	15.50–30.77
DMSO vs. TH ⁺	17.55	15.99–24.94
TH ⁻ vs. TH ⁺	19.13	15.92–27.90
AB pramaxilla development vs. Danionini premaxillae diversity	33.34	25.76–64.45
PC1-PC2		
AB vs. DMSO	61.51	50.06–85.52
AB vs. TH ⁻	88.02	46.32–94.51
AB vs. TH ⁺	86.73	47.20–91.12
DMSO vs. TH ⁻	65.94	40.42–90.94
DMSO vs. TH ⁺	74.15	42.80–91.19
TH ⁻ vs. TH ⁺	29.86	34.57–89.13
AB pramaxilla development vs. Danionini premaxillae diversity	52.15	38.73–84.78
PC1-PC3		
AB vs. DMSO	80.03	73.32–96.61
AB vs. TH ⁻	65.80	55.39–102.35
AB vs. TH ⁺	59.58	53.65–86.61
DMSO vs. TH ⁻	84.21	67.73–102.18
DMSO vs. TH ⁺	77.30	65.46–100.39
AB pramaxilla development vs. Danionini premaxillae diversity	52.83	50.17–96.84
PC1-PC4		
AB vs. DMSO	101.83	84.63–116.01
AB vs. TH ⁻	93.18	71.75–109.74
AB vs. TH ⁺	77.48	66.79–106.49
DMSO vs. TH ⁻	86.38	74.52–111.07
DMSO vs. TH ⁺	97.36	77.98–117.97
AB pramaxilla development vs. Danionini premaxillae diversity	79.00	74.07–109.85

Note: The observed angle between the shape spaces is given for each comparison. 95% confidence intervals (CIs) for this angle were calculated by bootstrapping the data from both groups (700 bootstraps). Significant differences in bold/shaded. The TH⁻ and TH⁺ shape spaces were not significantly different for the first PC axis only. All other comparisons were not significantly different for PC1-PC4.