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One size does not fit all: Variation in anatomical traits associated with emersion behavior in mudskippers (Gobiidae: Oxudercinae)

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Life histories involving transitions between differing habitats (i.e., aquatic to terrestrial or marine to freshwater) require numerous anatomical, physiological, and behavioral changes. Often, the traits associated with these changes are thought to come in suites, but all traits thought to be associated with particular life histories may not be required. While some traits are found in all species with a particular habitat transition, a grab bag approach may apply to other traits in that any trait may be sufficient for successful habitat transitions. We examine patterns of morphological traits associated with prolonged emersion in mudskipper, an amphibious fishes clade, where prolonged emersion appears twice. We test the evolutionary history of multiple characteristics associated with cutaneous respiration. We find most traits thought to be key for prolonged emersion show no phylogenetic signal and no tight correlation with prolonged emersion. Such traits appear in species with prolonged emersion but also non-emerging species. Only capillary density, which, when increased, allows for increased oxygen absorption, shows strong phylogenetic signal and correlation with prolonged emersion. Further experimental, functional genomics, and observational studies are needed to fully understand the mechanisms associated with each of these traits. With respect to traits associated with other particular behaviors, a comparative framework can be helpful in identifying evolutionary correlates.

KEYWORDS

amphibious, fish out of water, goby, histology, Pagel's lambda, phylogenetic comparative methods, trait evolution, water-to-land transition

Introduction

The ability to survive in a new environment often requires numerous behavioral, anatomical, and physiological traits. Life histories that allow for movement between habitats, such as transitions between marine and freshwater, aquatic and terrestrial, and flight and flightless, are all common across the tree of life (Shubin et al., 2006; Corush, 2019;

Sackton et al., 2019). In some cases, the traits associated with these life histories, or simply occupying a new environment in the case of migration or invasion, are thought to come in suites (Dingle, 2006; Colmer and Voesenek, 2009; Carnicer et al., 2013; Chuang and Riechert, 2021; but also see Piersma et al., 2005). It is convenient to consider the full suite of traits associated with each life history to be necessary. But parsing apart the traits that are necessary for persistence, as opposed to additional adaptive traits that are not necessary for survival in a new environment, is important in order to understand the evolution of such life histories. This is especially problematic in a single-species framework where a group of traits is associated with a life history and then each trait is assumed to be necessary. Examining suites of traits in a phylogenetic framework can be more useful in identifying specific traits that may be required for each life history.

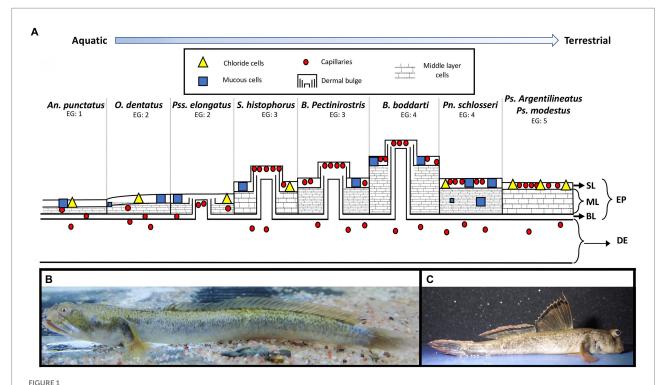
Traits associated with aquatic-terrestrial movement in fishes have been examined with respect to fin kinematics (Pace and Gibb, 2009), morphological and molecular basis of vision (Sayer, 2005; Hu et al., 2022), vasculatures of skin, gills, and buccoopercular cavities associated with respiration (Zhang et al., 2000; Gonzales et al., 2011), body elongation and compressions (Polgar et al., 2017; Egan et al., 2021), specialization in feeding and diet (Kruitwagen et al., 2007; Ord and Hundt, 2020; Tran et al., 2022), shifts in dominant bacteria in the gut (Guan et al., 2021), vocalizations (Polgar et al., 2011), burrowing behaviors (Ishimatsu et al., 1998; Dinh et al., 2021), candidate genes associated with emersion (You et al., 2014), and the effects of amphibious behavior on population structure (Corush et al., 2022). In many of these examples, multiple behavioral, physiological, and anatomical traits work together to allow fish to emerge. While few studies do take a comparative approach, e.g., two studies looking at body shape and diet of blennies (Blenniidae) in supralittoral and intertidal zones (Ord and Hundt, 2020; Egan et al., 2021), these traits are typically examined in a single, if not a few, species, and also in species that show a more extreme level of transition, making it tempting to assume that the suite of traits are all required for the evolution of the behavior. Speciose groups such as mudskippers (Gobiidae: Oxudercidae: subfamily Oxudercinae), however, have a complex evolutionary history with respect to the polyphyletic distribution of prolonged emersion (Steppan et al., 2022). This raises questions regarding the evolutionary history of the various traits associated with emersion. Understanding the phylogenetic signal of these traits and their correlation to emersion behavior is important for a broader understanding of how and why fish move

Respiration in emerging fishes is of interest because most fish species are unable to maintain necessary oxygen levels out of water. Across fishes that can utilize atmospheric oxygen, a number of organs are used including gills, skin, gastrointestinal tract, labyrinthine organs, and the swim bladder [for an in-depth review of air breathing and gas exchange in fishes see Graham (1997); Sayer (2005); Ishimatsu (2017) and Zaccone et al. (2018)]. Mudskippers, however, lack organs such as the gas bladder (Polgar et al., 2011) which is used by other air-breathing fishes including

gars (*Lepisosteus* and *Atractosteus*). The skin is important in mudskippers as cutaneous respiration is a primary respiration method for some species when out of water, although gills are also used (Tamura et al., 1976) as well as bucco-opercular cavity (Ishimatsu, 2017). Within mudskippers, there is variation in gas exchange. For example, *Periophthalmodon schlosseri* are more effective in oxygen uptake out of water compared to in water (Kok et al., 1998) even after "exhaustive exercise" (Takeda et al., 1999). Others, such as *Boleophthalmus boddaerti*, rely more heavily on aquatic respiration, although they are capable of surviving long periods out of water (Kok et al., 1998). Other species within the mudskipper clade are fully aquatic, although some such as *Apocryptodon punctatus* may gulp air as suggested by Graham and Lee (2004).

One organ that has received particularly wide taxonomical attention is the skin (Al-Kadhomiy and Hughes, 1988; Yokoya and Tamura, 1992; Zhang et al., 2000; Park, 2002; Park et al., 2003; Zhang et al., 2003; Park et al., 2006; Beon et al., 2013; Jaafar and Parenti, 2017; Kim and Park, 2020). Within the dermal layers, there is a suite of traits that increases the ability to absorb oxygen when fishes are out of water. These traits include an increase in capillary density in the skin allowing for more absorption, as well as a decrease in the distance between those capillaries and the surface of the skin, and decrease in the distance oxygen must travel to get to the blood vessels (Al-Kadhomiy and Hughes, 1988; Beon et al., 2013). Additionally, dermal bulges often contain additional capillaries and are associated with cutaneous respiration (Beon et al., 2013). The presence of mucous cells has been observed to decrease within species during emersion (Sayer and Blackstock, 1988) as the mucus may restrict oxygen flow (Zhang, 2001) although it is also suggested that this may help increase diffusion (Graham, 1997). Chloride cells are associated with ion regulation and are often associated with respiratory organs (Copeland, 1948; Komnick, 1986). Middle cells may also act as a barrier to water loss and store large amounts of water (Yokoya and Tamura, 1992; Figure 1). Because most of these traits do not require a novel feature, but rather an increase or decrease in an already existent trait (i.e., more blood capillary or thicker cells), the "gain" (increase in the trait) and "loss" (decrease in the trait) is likely more reversible as opposed to traits typically thought of in the context of Dollo's law (Dollo, 1893) which states that once a complex trait is lost, it should not be reacquired (e.g., the regain of a digit in the lizard genus Bachia: Galis et al., 2010).

Of the seven genera of mudskippers, there is a wide range of emersion behavior from fully aquatic, emersion group (EG) 1, to primarily terrestrial, EG 5 (See Supplementary Table 1 for the description of each EG). According to recent phylogenetic analyses (McCraney et al., 2020; Steppan et al., 2022), there are two monophyletic groupings containing the species known for prolonged emersion (EG 4–5). The first clade contains the species of the genera *Periophthalmus* and *Periophthalmodon* (clade P: EG 4–5) and the second clade, nested within a less frequent, less prolonged emersion species, is the genus *Boleophthalmus* (clade B: EG 4, but also contains species in EG 3; Figure 2). It is unknown



(A) Schematic representation of the variation in histological features from the head of mudskipper species representing different emission groups from fully aquatic (emersion group EG1) (left) to highly terrestrial (EG5) (Right). Layers include the dermis (DE) and epidermis (EP), which included a superficial layer (SL), middle layer (ML), and basal layer (BL). Figure modified from Zhang (2001). Morphological variation between EGs can be seen between (B), Oxuderces dentatus (EG2) collected in Panyu, Guangdong Province, China, in September 2022 (Photo credit: Minzheng Li) and (C), Periophthalmus magnuspinnatus (EG5) (not used in this study, but representative of the "P clade" in figure 2) collected from the Bohai Sea, China, in July 2016 (Photo credit: Joel B. Corush).

how the traits associated with prolonged emersion have changed across the phylogeny. Additionally, it is unknown if the same suite of traits is found in all species with prolonged emersion. If only a few adaptive traits are needed to successfully spend time out of water, different species might exhibit different traits. These points can be addressed by understanding the phylogenetic signal of traits associated with emersion.

Phylogenetic signal is a measurement of the randomness of a trait's distribution across a phylogeny. If a trait is randomly distributed across a phylogeny, then there is no signal. If this is the case, traits are not often shared by sister species or closely related species, but instead found randomly across the phylogeny. However, if a trait shows a nonrandom pattern with respect to phylogeny, then we could consider there to be a strong phylogenetic signal. In this case, traits tend to be found in closely related species or conversely, not found in closely related species. Two standard metrics of phylogenetic signal include Pagel's λ (Pagel, 1997; Pagel, 1999) and Blomberg's K (Blomberg et al., 2003). Pagel's λ ranges from 0 (no phylogenetic signal) to 1 (strong phylogenetic signal). Blomberg's K starts from 0 (no phylogenetic signal) and continues infinitely with K=1 being a threshold for phylogenetic signal and increasing values indicating a stronger signal. Strength of phylogenetic signal alone is not an indicator of an association between traits. Two traits may both have strong phylogenetic signals because they are restricted to different clades.

For example, in this study, we found a strong phylogenetic signal for dermal bulges and emersion behavior, but these traits appear in different clades. Additional tests are needed to show the association between traits.

We used histological data from Zhang (2001), coupled with a published phylogeny (McCraney et al., 2020) to identify the evolutionary pattern of skin adaptations across multiple species of mudskippers. This suite of traits was used in part because of the availability of data on multiple traits across species that do and do not emerge. We address the following questions: (1) Are the specific traits that are thought to be associated with prolonged emersion present in all species with prolonged emersion and absent in species without prolonged emersion? (2) Do the traits appear in suites or are these traits irregularly distributed across species with prolonged emersion? (3) Is there a phylogenetic signal associated with each of the traits?

Materials and methods

Histology

Histological data were summarized by Zhang (2001) for the giant mudskipper (*Periophthalmodon schlosseri*; Pallas, 1774), the barred mudskipper (*Periophthalmus argentilineatus*; Cuvier and

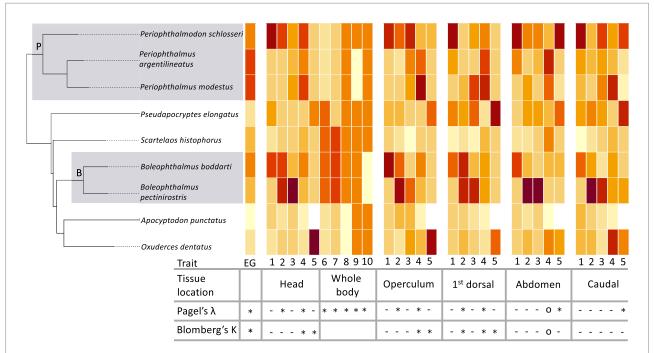
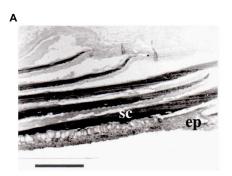


FIGURE 2
Pruned phylogeny (from McCraney et al., 2020) and a heatmap of associated trait values. Traits estimation based on mean values of each species which is based on mean values for each. Traits include: (1) number of scale layers (#), (2) tier of middle cells (#), (3) middle layer thickness (μ m), (4) capillary density (#/mm), (5) diffusion distance (μ m), (6) dermal bulge, (7) papilla, (8) Intra-epidermal capillary (skin), (9) mucous cells, and (10) chloride cell. Traits 6–10 are presence/absence across the full specimen. Dark colors=higher/presence. Light colors=low/absence trait values (see Supplementary Table 2 for actual values). Shaded boxes indicate clade P and clade B which have prolonged emersion. Table below represents phylogenetic signal based on Pagel's λ and Blomberg's K (See supplementary Table 4 for actual values). Traits and tissues combinations that show a strong phylogenetic signal are denoted with an "** (λ >0.9, K>1.0), traits and tissues combinations that show a little to no phylogenetic signal are denoted with an "o" (0.5< λ <0.9, 0.9<K<1.0), and traits and tissues combinations that show a little to no phylogenetic signal are denoted with a "-" (λ <0.5, K<0.9). Blomberg's K was not calculated for the whole body binary traits.

Valenciennes, 1837), shuttles hoppfish (Periophthalmus modestus; Cantor, 1842), the elongate mudskipper (Pseudapocryptes elongatus; Cuvier, 1816), the walking goby (Scartelaos histophorus; Cuvier and Valenciennes, 1837), Boddart's goggle-eyed goby (Boleophthalmus boddarti; Pallas, 1774), the great blue-spotted mudskipper (Boleophthalmus pectinirostris; Linnaeus, 1758), the spotted hidden-teeth goby (Apocryptodon punctatus; Tomiyama, 1934), and the crocodile-face goby (Oxuderces dentatus; Eydoux and Souleyet, 1852). For each sample, measurements were taken from the following regions of the body: (1) top of the head (H), (2) outer surface of operculum (O), (3) dorso-lateral body near the first dorsal fin (D), (4) posterior portion of the abdomen (A), and (5) the lateral side of the caudal peduncle (C). Traits measured at each of the regions included: (1) number of scale layers (#), (2) tier of middle cells (#), (3) middle layer thickness (µm), (4) capillary density (#/mm), and (5) diffusion distance (µm). Additionally, the following traits were examined as presence/absence across the full specimen: (6) drmal bulge, (7) papilla, (8) intra-epidermal capillary, (9) mucous cells, and (10) chloride cell. Lastly, each species was assigned an emersion period (EG 1-5) based on how much time they spent out of water. Values used for phylogenetic analyses were the means across all individuals of each species as recorded in Zhang (2001). If minimum and maximum values were measured in Zhang (2001), as in tier of middle cells and middle layer thickness, the mean of the minimum and maximum was used (Supplementary Table 2). The number of scale layers was considered 0 if there was not a complete layer of scales (i.e., noncontinuous scales that are completely embedded in the skin with spacing between scales; Figure 3). A heatmap of trait values was constructed using the 'heatmap' function in the basic stats package in R (R Core Team, 2019).

Phylogenetic analysis

A recent goby phylogeny (McCraney et al., 2020) was pruned to include only the taxa of interest (Figure 2) using the "drop.tip" function in the ape package (Paradis et al., 2004). Phylogenetic signal was calculated using the "phylosig" function in the phytools package with both method = "K" and "lambda" (Revell, 2012). For discrete traits, the "fitDiscrete" function in Geiger (Harmon et al., 2008) was used to calculate Pagel's λ . Because of the small number of taxa in this study, a permutation test was performed to assess the significance of these values. The permutation test was performed by randomly arranging the data and recalculating both metrics of phylogenetic signal. This was done 1,000 times for each



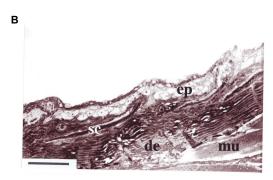


FIGURE 3

Examples of variation in scale counts from different species of mudskipper from Zhang (2001). Images depict the (A), transverse section of the operculum of *Periophthalmodon septemradiatus* (not included in this study) and (B), lateral side of the caudal peduncle of *Scartelaos histophorus*. sc, scales; ep, epidermis; de, dermis; mu, muscle. Scale bars indicate 100 μm. *Pn. septemradiatus* has continuous multilayered scales, whereas *S. histophorus* has isolated embedded scales.

trait and tissue type using Pagel's λ and Blomberg's K. Observed values were compared to the distribution of simulated values.

To identify the correlation between traits, Pearson's productmoment correlation coefficient (r) was calculated using the "cor. test" with methods = "pearson" in the basic stats package in R. In order to incorporate evolutionary history, two separate methods were used depending on the type of data. To look at correlations, phylogenetic generalized least squares (PGLS) were used to identify significant associations between pairs of traits. This was done using the "glm" function in the nlem package (Pinheiro et al., 2017). For the correlation argument, phylogeny was incorporated with the "corBrownian" function in the ape package. Because PGLS should not be used between two binary traits, phylogenetic generalized linear mixed models (PGLMM) were used to identify significant associations between pairs of binary traits (Ives and Garland, 2014; Symonds and Blomberg, 2014). To look at the correlation between binary traits (all the non-tissue specific traits dermal bulge, papilla, intra-epidermal capillary, mucous cells, and chloride cells), the "binaryPGLMM" function in the ape package (Pearse et al., 2015) was used. Because of the number of independent correlations, a Bonferroni correction for multiple comparisons was completed using the "p.adjust" function in the basic stats function in R (see supplementary file for example code).

Results/discussion

We find that while some of the traits associated with prolonged emersion do tend to show up in all species with prolonged emersion and not others, this is not the observed pattern across most traits. Many traits thought to be associated with emersion are randomly distributed in species in both high and low EGs. This is evident by the β coefficient of only one trait, capillary density, showing a significant effect of emersion groups when incorporating phylogenetic history (Supplementary Table 3). This suggests that increased capillary density may be a key component of emersion in mudskippers. It is also worth noting that this trait showed a

significant correlation in tissues H (β =0.109, p=0.011), O (β =0.137, p=0.026), D (β =0.109, p=0.005), but not A (β =0.349, p=1.0), and C (β =0.167, p=1.0) after corrections for multiple comparisons. This is biologically meaningful because even species such as *Periophthalmus modestus* and *Periophthalmus argentilineatus*, which are both in EG 5, often have their abdominal region and their tails in, or pressed against, water as they sit on wet mud, or on a rock or log with the posterior part of their body submerged. Because those body parts are not being exposed to air, individuals may not allocate resources to increase capillary density in those regions.

Traits associated with prolonged emersion do not appear to show up in suites and very few traits showed any correlation with other traits. Two traits that did show near significant correlation in tissues A (β = 0.034, p = 0.055) but not C (β = 0.083, p = 1.0), D $(\beta = 0.082, p = 1.0)$, H $(\beta = 0.054, p = 0.22)$, or tissues O $(\beta = 0.062, p = 0.062)$ p=1.0; Supplementary Table 3), were tier of middle cells and middle layer thickness, which would be expected since an increased number of cells in a layer should increase the overall layer thickness. Additionally, tier of middle cells shows a significant association with chloride cells, but only in tissue D $(\beta = -9.545, p < 0.001)$. The importance of incorporating phylogenetic independent contrasts can be seen by the numerous traits that did show a significant r, but not when incorporating the phylogenetic relationship. The overall lack of correlation in traits implies that, while there may be various traits that are adaptive with respect to emersion behavior, different lineages may make use of different combinations of traits (Supplementary Table 3).

Lastly, many of the traits do appear to show a phylogenetic signal (Figure 2). Emersion group did show a strong phylogenetic signal with both Pagel's λ (λ =0.999) and Blomberg's K (K=1.608) as did dermal bulge, papilla, mucous cells, intra-epidermal capillary (skin), and chloride cell. Some traits, such as dermal bulges, have a very apparent phylogenetic signal, but are found in the species with the highest EG (e.g., Clade B) as well as species with the lowest EG (e.g., *Pseudapocryptes elongatus*). Capillary density also showed a strong phylogenetic signal according to both metrics in tissues H, O, and D, but not in A or C (Figure 2,

Supplementary Table 4). This is also the only trait that shows a strong correlation with EG. The permutation tests of all traits showing a strong phylogenetic signal resulted in observed values of Pagel's λ and Blomberg's K to be significantly higher than randomized data (p < 0.05; Supplementary Figure 1).

Both phylogenetic and histological data are subject to some level of variation based on the methods used. We acknowledge that scales may be lost during handling of specimens, but the lack of scales on the operculum of many species is consistent with Boleophthalmus species' descriptions by Murdy (1989). The lack of scales on the head of Oxuderces dentatus is also consistent with previous descriptions (Jaafar and Parenti, 2017). Additionally, Murdy (1989) describes a widespread lack of scales on Zappa, while not a represented genus in our study, this suggests there is variability within the clade. Therefore, the lack of scales in the Scartelaos is assumed to be correct, not a product of handling the specimens. A larger number of samples would be helpful in a better understanding of the species-specific variation found in these traits. This could be particularly important in species that are found in both temperate and tropical regions (e.g., Periophthalmus argentilineatus) as they may alter their morphology to adjust to local conditions. Additional data on the remaining taxa in the clade, particularly unsampled genera (e.g., Apocryptes, Parapocryptes, Zappa, Taenioides, Odontamblyopus) as well as a more complete phylogeny will better inform our understanding of how these adaptive traits have evolved across the Oxudercinae radiation. The increase in annotated mudskipper genomes (Li et al., 2022; Yang et al., 2022) may also help identify specific candidate genes associated with specific traits. Furthermore, the location of the sampled skin is important to consider (Zhang et al., 2003), species such as those in the genus Oxuderces may not fully emerge, with the ventral portion of their body remaining submerged all the time. This should result in the skin on their head having traits more closely associated with emersion whereas the skin on the ventral side has traits more closely related with a typical fish.

These results can be helpful in identifying specific adaptive traits worth further examination when addressing questions about emersion behavior. If the patterns observed in this study are in fact causal, we expect to see the change in capillary density expanded across the full mudskipper clade. Similar mechanisms may also be observed across other groups that emerge, such as blennies and catfishes, although these groups may utilize other traits. This is also interesting in the context of individual and species-specific variation. Mudskippers have a fully aquatic larval phase which may require less vascularization. Additionally, humidity and temperature differences in temperate compared to tropical environments may lead to different mechanisms in oxygen uptake in species that span both environments such as Periophthalmus argentilineatus. This methodology can also be applied to identify behavioral, anatomical, and physiological correlates to other life histories involving transitions between habitats such as marine to freshwater, shifts in thermal tolerance, invasive species, or species' response to climate change.

Data availability statement

The original contributions presented in the study are included in the article/Supplementary material, further inquiries can be directed to the corresponding authors.

Author contributions

JC and JZ designed the study. JZ collected the histological data as part of her dissertation research (Zhang, 2001). JC conducted the phylogenetic analysis. JC prepared the manuscript with input from JZ. All authors contributed to the article and approved the submitted version.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Supplementary material

The Supplementary material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fevo.2022.967067/full#supplementary-material

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