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2	Testing the behavioral origins of novelty: did increased
3	aggression lead to scale-eating in pupfishes?
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23 Abstract

24 How novelty evolves is still largely unknown. Environmental changes are often assumed to 25 precede novelty; however, behavioral shifts may also play a role. Here, we examine whether a 26 shift in aggression explains the origin of a novel scale-eating pupfish species (Cyprinodon 27 desquamator) within an adaptive radiation on San Salvador Island, Bahamas. We compared 28 aggression using behavioral and gene expression data across three sympatric species in the San 29 Salvador radiation (generalist, snail-eating specialist, and scale-eating specialist), and 30 additionally measured behavioral aggression in an outgroup generalist from North Carolina. 31 Surprisingly, we found increased behavioral aggression and differential expression of 32 aggression-related genes in both the scale-eating and snail-eating species. Furthermore, male 33 scale-eaters and female snail-eaters showed the highest levels of aggression compared to other 34 groups. Differential gene expression in each specialist during larval development also suggested 35 sex-mediated differences in male-male aggression and maternal care. Ultimately, our data 36 indicate that aggression is not unique to scale-eating specialists. Instead, selection may increase 37 aggression in other contexts such as niche specialization, mate competition, or selection on other 38 ecologically relevant traits, including jaw size. Indeed, some adaptive variants associated with 39 oral jaw size in the San Salvador radiation occur in genetic pathways with pleiotropic effects on 40 aggression.

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47 Introduction

48 The origins of evolutionary novelty are still poorly understood. For example, both novel 49 behavior and morphology play a role in the evolution of novel resource use, but their relative 50 importance and order in which they evolve are still unknown. Changes in behavior may precede 51 the evolution of novel morphologies, as they can expose organisms to novel environments and 52 selective pressures (Huey et al. 2003). Investigations of novelty, however, overwhelmingly 53 ignore this possibility (although see: Huey et al. 2003; Losos et al. 2004; Duckworth 2006). 54 Instead, previous studies have focused on novel adaptive morphologies or on how environmental 55 changes expose organisms to new selective pressures (Liem 1973; Barton and Partridge 2000; 56 Janovetz 2005a; Hulsey et al. 2008). One reason behavior has been overlooked as an origin for 57 novelty is because it is still unclear whether it drives or inhibits evolution, as behavior is often 58 extremely plastic. However, in order to determine if novelty has a behavioral origin we must 59 first understand its variation within and among taxa. 60 One outstanding example of novelty is lepidophagy (scale-eating) in fishes. Scale-eating 61 has been documented in five freshwater and seven saltwater families of fishes and has 62 independently evolved at least 19 times (Sazima 1983; Janovetz 2005b; Martin and Feinstein 63 2014; Kolmann et al. 2018). Scale-eating includes both novel morphologies and novel behaviors. 64 For example, some scale-eaters have premaxillary external teeth for scraping scales 65 (Novakowski et al. 2004), some use aggressive mimicry to secure their prey (Boileau et al. 66 2015), others sneak scales from the surface of fish that they are cleaning (Losey 1979), and still 67 others use ambush tactics to obtain scales (Nshombo et al. 1985). Even though scale-eating is an 68 outstanding example of the convergent evolution of novel trophic ecology across disparate

69 environments and taxa and displays a wide variety of morphologies and behaviors, its origins70 have yet to be explored.

71 There are currently only three hypothesized behavioral origins for scale-eating. The 72 algae-grazer hypothesis predicts that scale-eating arises from the incidental ingestion of scales 73 during algae scraping (Fryer et al. 1955; Greenwood 1965; Sazima 1983). Many scale-eaters are 74 closely related to algae-grazers. For example, many rock-dwelling Malawi cichlids are algae-75 scrapers (Greenwood 1965; Fryer and Iles 1972; Ribbink et al. 1983), but the radiation also 76 includes two sister species of scale-eaters (*Corematodus shiranus* and *Corematodus taeniatus*) 77 and a second independent origin of scale-eating in *Genvochromis mento* (Trewavas 1947; 78 Greenwood 1965). Similarly, the extinct Lake Victorian scale-eater *Haplochromis welcommei* 79 was nested within rock-dwelling algae scrapers (Greenwood 1965). This hypothesis, however, 80 does not address why algae-grazing fish would seek food on the surface of other fish 81 (Greenwood 1965). The second hypothesis, termed the cleaner hypothesis, tries to address this 82 gap by arguing that scale-eating arose from the incidental ingestion of scales during the 83 consumption of ectoparasites from the surface of other fishes (Greenwood 1965; Sazima 1983). 84 One line of evidence supporting this hypothesis is that cleaner fish sometimes eat scales. For 85 example, the Hawaiian cleaner wrasse (*Labroides phthirophagus*) and two species of juvenile 86 sea chub (Hermosilla azurea and Girella nigricans) consume both ectoparasites and scales 87 (Demartini and Cover 1981; Sazima 1983; Losey 1972). However, most scale-eating fishes are 88 not known to forage on ectoparasites, nor are they closely related to fish that do. In fact, the 89 closest examples of this are the false cleaner fishes (Aspidontus taeniatus and Plagiotremus 90 rhinorhynchs) who aggressively mimic cleaner wrasse (Labroides dimidiatus) in order to 91 consume scales. Despite their morphological similarities, however, these fish are not closely

92 related (Hundt et al. 2014). Finally, the aggression hypothesis predicts that scale-eating evolved 93 due to the accidental ingestion of scales during inter- or intra-species aggression (Sazima 1983). 94 This hypothesis is supported by the fact that two characid species of scale-eaters (*Probolodus* 95 *heterostomus* and *Exodon paradoxus*) are closely related to the aggressive *Astyanax* tetras 96 (Sazima 1983; Kolmann et al. 2018); a similar argument can be made for the scale-eating 97 piranha (Cataprion mento) (Janovetz 2005). 98 The scale-eating pupfish, *Cyprinodon desquamator*, is an excellent species for 99 investigating the origins of scale-eating because it is, by far, the youngest scale-eating specialist. 100 The species is nested within a sympatric adaptive radiation of pupfishes endemic to the 101 hypersaline lakes of San Salvador island, Bahamas (Martin and Wainwright 2011, 2013a). In 102 addition to the scale-eating pupfish, this radiation also includes a widespread generalist (C. 103 *variegatus*) and an endemic snail-eating specialist (*C. brontotheroides*). Other generalist pupfish 104 lineages are also distributed across the Caribbean and western Atlantic Ocean. Phylogenetic 105 evidence indicates that scale-eater's most recent ancestor was a generalist feeder (Martin 2016; 106 Richards and Martin 2017). Furthermore, geological evidence suggests that the hypersaline 107 lakes of San Salvador island—and thus the radiation containing the scale-eater—is less than 10 108 thousand years old (Hagey and Mylroie 1995; Martin and Wainwright 2013a,b). 109 We investigated the possible behavioral origins of novelty by examining whether a shift 110 in aggression led to the evolution of scale-eating in pupfish. We compared measures of 111 aggression using both behavioral and gene expression data between scale-eaters and closely 112 related species within their radiation (a sympatric generalist species and a snail-eating species 113 from San Salvador Island) as well as an additional outgroup (a generalist species from North

114 Carolina). We predicted high levels of aggression in scale-eaters, intermediate levels in the San

115	Salvador generalist and snail-eating pupfish, and low levels of aggression in the North Carolina
116	(NC) outgroup generalist. Surprisingly, we found that male scale-eaters and female snail-eaters
117	displayed increased levels of aggression, while other groups displayed relatively low levels of
118	aggression. This suggests that aggression alone cannot explain the origins of scale-eating in
119	pupfish. We also identified promising candidate genes that are differentially expressed between
120	species associated with differences in aggression and linking behavior, morphology, and the
121	evolution of novel ecology.
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123	Methods
124	Sampling
125	Generalist, snail-eating, and scale-eating pupfish were collected by seine from Crescent Pond,
126	Great Lake, Little Lake, Osprey Lake, and Oyster Pond of San Salvador Island, Bahamas in July,
127	2016. In June 2017, generalist pupfish were collected by seine from the Cape Fear river drainage
128	(Fort Fisher) on the coast of North Carolina. Fishes were housed in $40 - 80$ liter tanks in mixed-
129	sex groups at 5-10 ppt salinity in temperatures ranging from 23°C-30°C. Fish were fed a diet of
130	frozen blood worms, frozen mysis shrimp, or commercial pellet food daily.
131	
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133 We quantified levels of aggression for each pupfish species and sex using mirror tests (Vøllestad

134 and Quinn 2003; Francis 2010). To control for individual size and motivation, we incited

- aggression using a compact mirror (10 cm X 14 cm) placed in a 2-liter trial tank (25 cm X 16 cm
- 136 X 17 cm). We randomly chose adult fish and isolated each one in 2-liter tanks that contained a

137	single bottom synthetic yarn mop for cover and opaque barriers between adjacent tanks. We gave
138	the fish at least 12 hours to acclimate to their new environment before performing an assay.
139	During a 5-minute focal observation period, we measured three metrics as a proxy for
140	aggression: latency to approach mirror image, latency to attack mirror image, and total number
141	of attacks toward the mirror image. A trial began as soon as the mirror was securely lowered into
142	the tank. We measured latency to approach as the time elapsed before an individual approached
143	the mirror to within one-body length. Similarly, we measured latency to attack as the time
144	elapsed before an individual attacked their mirror image for the first time. Finally, we counted
145	the total number of attacks an individual performed during the entirety of the trial. We also
146	measured the standard length of each fish after the trial.
147 148 149	Statistical analyses
150	We used time-to-event analyses to determine if species and sex were associated with 1) latency
151	to approach their mirror image and 2) latency to attack their mirror image. For the latency to
152	approach metric (time in seconds) and the latency to attack metric (time in seconds) we used a
153	mixed-effects Cox proportional hazards model (coxme package; Therneau 2012) in R (R
154	Development Core Team 2016). These models allow right censored data, i.e. individuals who did
155	not approach or attack their mirror image were not excluded and contributed to Kaplan-Meier
156	
	estimates and time-to-event curves (Rich et al. 2010). For both the latency to approach model
157	estimates and time-to-event curves (Rich et al. 2010). For both the latency to approach model and the latency to attack model we included species, sex, and their interaction as fixed effects
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	and the latency to attack model we included species, sex, and their interaction as fixed effects
158	and the latency to attack model we included species, sex, and their interaction as fixed effects and lake population as a random effect. We compared these models to equivalent models that

162 however, size was a significant covariate and retained in the final model. We used the likelihood 163 ratio test to determine if species, sex, or their interaction were associated with latency to 164 approach or attack the mirror image. Additionally, we used a Cox proportional hazards model 165 without mixed effects to plot the resulting time-to-event curves and made pairwise comparisons 166 between curves using log-rank tests (Survival Package; survminer package; Therneau 2015). 167 We analyzed the total number of attacks using a generalized linear mixed model 168 (GLMM) with a negative binomial distribution for this response variable. We modeled species, 169 sex, and their interaction as fixed effects, and population as a random effect. We compared this 170 model to a model including size (log scale) as a continuous covariate using AICc, but accounting 171 for size did not substantially increase the likelihood of the model ($\Delta AICc = 0.96$). We used a 172 Wald chi-square test (type II) to determine if species, sex, or their interaction significantly 173 affected the total number of attacks performed and used Tukey's HSD to make direct 174 comparisons between groups.

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176 Identifying candidate genes affecting differences in aggression between species

177 We searched a previously published dataset of 15 San Salvador pupfish transcriptomes to 178 identify candidate genes underlying behavioral differences in aggression among all three species 179 (Mcgirr and Martin 2018). This previous study did not analyze gene expression pathways 180 annotated for effects on behavior. Briefly, purebred F_1 and F_2 offspring from the three-species 181 found on San Salvador island were raised in a common garden laboratory environment. Larvae 182 were euthanized in an overdose of MS-222 at 8-10 days post fertilization (dpf) and were 183 immediately preserved in RNAlater (Ambion, Inc.) and stored at -20 C after 24 hours at 4 C. 184 Total mRNA was extracted from 6 generalists, 6 snail-eaters, and 3 scale-eaters (RNeasy kits,

185	Qiagen). Stranded sequencing on an Illumina HiSeq 4000 at the High Throughput Genomic
186	Sequencing Facility at UNC Chapel Hill resulted in 363 million raw reads that were aligned to
187	the Cyprinodon variegatus reference genome (NCBI, C. variegatus Annotation Release 100,
188	total sequence length =1,035,184,475; number of scaffold=9,259, scaffold N50, =835,301; contig
189	N50=20,803; Lencer et al. 2017). Aligned reads were mapped to annotated features using STAR
190	(v. 2.5(33)), with an average read depth of 309x per individual and read counts were generated
191	across annotated features using the featureCounts function from the Rsubread package (Liao et
192	al. 2013). DEseq2 (Love et al. 2014, v. 3.5) was used to normalize counts and identify: 1) 1,014
193	differentially expressed genes between snail-eaters vs generalists and 2) 5,982 differentially
194	expressed genes between scale-eaters vs generalists (McGirr and Martin 2018).
195	We identified one-way best hit zebrafish orthologs for genes differentially expressed
196	between 1) snail-eaters vs generalists (n=722) and 2) scale-eaters vs generalists (n=3,966) using
197	BlastP (V. 2.6; E-value <1). We compared this list of orthologs to gene ontologies describing
198	aggressive behavior (GO: 0002118), inter-male aggressive behavior (GO: 0002121), maternal
199	aggressive behavior (GO:0002125), maternal care behavior (GO: 0042711), and territorial
200	aggressive behavior (GO: 0002124; AmiGo; Carbon et al. 2009; Ashburner et al. 2000;
201	The Gene Ontology Consortium 2017). We also searched gene ontologies for three hormone
202	pathways commonly associated with aggression (the vasopressin pathway, the androgen
203	pathway, and the estradiol pathway).
204	

205 **Results**

206 Behavioral aggression

207 Male scale-eaters and female snail-eaters showed increased levels of aggression compared to 208 other groups. Scale-eaters (at the species level) approached their mirror image significantly more 209 than NC generalists (Table 1A; Fig. 1A; log-rank test, P = .038). Additionally, male scale-eaters 210 attacked their mirror image significantly more than male San Salvador generalists (Table 1B; 211 Fig. 1B; log-rank test, P = 0.032). Lastly, male scale-eaters were the only group to exhibit 212 increased aggression compared to their female counterparts. More male scale-eaters attacked 213 their mirror image than did female scale-eaters (Table 1B; Fig. 1B: log-rank test, P = 0.003), and 214 they performed significantly more total attacks (Table 1C; Fig. 2; Tukey HSD, P = 0.0003). 215 Female snail-eaters showed a similar pattern of increased aggression. Snail-eaters (at the 216 species level) approached their mirror image significantly more than NC generalists (Table 1A; 217 Fig. 1A; log-rank test, P = .038). Significantly more female snail-eaters attacked their mirror 218 image than all other female groups (Table 1B; Fig. 1B; log-rank test, NC generalist P = 0.032, 219 San Salvador generalist P = 0.0027, San Salvador scale-eater P = 0.0081). Finally, female snail-220 eaters performed significantly more attacks than female San Salvador scale-eaters and generalists 221 (Table 1C; Fig. 2; Tukey HSD, generalist P = 0.0032, scale-eater P = 0.0006).

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223 Gene Expression

We searched genes that were differentially expressed between scale-eaters *vs* generalists or between snail-eaters *vs* generalists for gene ontologies describing aggressive behavior, inter-male aggressive behavior, maternal aggressive behavior, maternal care behavior, the vasopressin hormone pathway, and the androgen hormone pathway (Table 2). Despite over one thousand differentially expressed genes at this developmental stage, only five genes were associated with these aggression-related ontologies in the snail-eater *vs* generalist comparison (Table 2A). Scaleeaters also exhibited differential expression of genes associated with inter-male aggression and
vasopressin when compared to their generalist sister species (Table 2B). Interestingly, one of
these genes (*gnaq*) contains a fixed variant in scale-eaters, which is known to function in
craniofacial development and shows signs of a hard selective sweep (McGirr and Martin 2017;
McGirr JA unpublished data). None of these ontologies were significantly over-represented in
either species comparison, which were instead enriched for cranial skeleton, metabolism, and
pigmentation terms (McGirr and Martin 2018).

237

238 Discussion

239 The origins of novelty have overwhelmingly been examined from a morphological perspective, 240 often ignoring behavior's potential role. However, shifts in behavior may also be a viable origin 241 for novelty. Unfortunately, only a few previous studies have directly investigated this possibility. 242 The origin of novelty in the Pacific field cricket (*Teleogryllus oceanicus*)—which exhibits a 243 novel silent morph—is one of the few examples of evolutionary novelty with a behavioral origin 244 (Zuk et al. 2006; Tinghitella and Zuk 2009; Bailey et al. 2010). Increased brain size in birds has 245 also been linked to behavioral shifts and novelty. Birds that display innovative feeding behaviors 246 have larger brains and are more successful at invading novel environments (Nicolakakis and 247 Lefebvre 2000; Sol and Lefebvre 2000; Overington et al. 2009). Likewise, the role of behavior in 248 evolutionary novelty has also been explored in western bluebirds (Sialia Mexicana; Duckworth 249 2006) and Anolis lizards (Losos et al. 2004, 2006). Despite the growing empirical evidence of 250 behavior's role in evolutionary innovation, a consensus has not yet been reached on whether 251 behavior ultimately drives or inhibits novelty. Furthermore, studies that investigate behavioral 252 origins of novelty rarely do so using both behavioral and genetic approaches. However, by

253	leveraging our gene expression data, we gained some mechanistic insight into the divergent
254	origins of increased behavioral aggression in each specialist species.

255 We tested whether increased aggression contributed to the origin of scale-eating in a 256 species of Caribbean pupfish using both behavioral and gene expression data. The aggression 257 hypothesis predicts that scale-eating arose due to increased inter- and intra- specific aggression 258 (Sazima 1983). Contrary to these predictions, both snail-eaters and scale-eaters showed increased 259 levels of aggression. Our gene expression data supported these findings, as both scale-eaters and 260 snail-eaters showed differential expression of genes involved in several aggression-related 261 pathways during larval development. Additionally, both scale-eaters and snail-eaters displayed 262 surprising differences in aggression between the sexes. While male scale-eaters showed 263 increased levels of aggression, female scale-eaters showed extremely low levels of aggression. 264 Conversely, female snail-eaters showed increased levels of aggression compared to females of 265 other species. These results suggest that the aggression hypothesis alone cannot explain the 266 evolution of scale-eating. Instead, selection may have favored increased levels of aggression in 267 other contexts, such as mate competition or trophic specialization in general. Increased levels of 268 aggression could have also arisen indirectly due to selection for other behaviors or traits, 269 including several genes involved in both aggression and craniofacial morphology (e.g. gnaq). 270 One caveat is that there is still discussion whether mirror tests accurately predict levels of 271 aggression in the field. Balzarini et al. (2014) argue that, while mirror tests are a valid method of 272 measuring aggression in some species, they are inappropriate for others. For example, some 273 species use lateral displays of aggression which primarily occur head to tail—a maneuver that is 274 impossible with a mirror image. Additional studies also indicate that mirror tests may not 275 accurately predict aggressive display frequency, duration, or orientation (Elwood et al. 2014;

Arnott et al. 2016). It is possible that our method of measuring aggression may have
underestimated aggression for scale-eating pupfish. This may be particularly true for female
scale-eaters. Our study primarily measured direct displays of aggression (i.e. attacks), however,
females often display aggression indirectly (Rosvall 2011; Stockley and Campbell 2013). Our
methods of measuring aggression, therefore, may have missed increased levels of aggression in
female scale-eaters while still detecting them in males.

282 A second caveat is that we compared differential gene expression in an early larval 283 developmental stage, 8-10 dpf, long before sexual maturity in this species. Thus, we are not 284 comparing adult differences in gene expression between the sexes in each species. Instead, by 285 examining early larval stages our gene expression analyses provide insight into species-specific 286 differences in aggression-related genetic pathways established during an early developmental timepoint. This has the advantage of defining structural developmental differences in each 287 288 species, rather than transient differences in transcription between adult male and female fish 289 sensitive to dominance status, reproductive state, and mood. Furthermore, we found surprising 290 congruence between our behavioral and transcriptomic data supporting the conclusions of 291 increased aggression in both San Salvador specialists due to different aggression-related genetic 292 pathways.

293

294 New hypotheses for varying levels of aggression between pupfish species

If increased levels of aggression are not associated with scale-eating, then what explains this variation between species? One possibility is that selection may have directly favored increased aggression in the context of dietary specialization. Aggression may be positively correlated with traits associated with specialization (Genner et al. 1999; Peiman and Robinson 2010; Blowes et

299 al. 2013), suggesting that specialists should show increased levels of aggression compared to 300 generalists. Existing evidence supports this as increased levels of aggression have been 301 documented in specialist butterfly fish (*Chaetodontids*; Blowes et al. 2013), specialist striped 302 surfperch (*Embiotoca lateralis*; Holbrook and Schmitt 1992), and have even been observed in 303 game-theoretic simulation models (Chubaty et al. 2014). A second possibility is that increased 304 aggression may be associated with colonizing a novel niche. Aggression is tightly correlated with 305 boldness in a phenomenon termed the aggressiveness-boldness syndrome (Sih et al. 2004). Many 306 studies have shown that increased boldness in species such as cane toads, mosquitofish, and 307 Trinidadian killifish leads to increased dispersal into novel habitats or niches (Fraser et al. 2001; 308 Rehage and Sih 2004; Gruber et al. 2017). This relationship indicates that increased aggression 309 may be an incidental effect of selection for increased boldness and occupation of a novel niche. 310 Our data supports either scenario, as we observed increased aggression in both San Salvador specialists. Neither of these hypotheses, however, explain the variation in aggression between 311 312 sexes.

313

314 Aggression and mating system

Increased aggression may be favored in the context of courtship or mate competition. It is well documented across multiple taxa that the sex with the higher potential reproductive rate should have increased levels of aggression as they must compete more heavily for access to mates (Clutton-Brock and Parker 1992; Andersson 1994; Jennions and Petrie 2007). Normally, males have higher potential reproductive rates since mating is energetically cheap for them (Trivers Scale-eaters, and *Cyprinodon* pupfishes in general, seem to adhere to this standard as they mate in a lekking system and do not provide parental care (Gumm 2012). Male scale-eaters may

322 be more aggressive to compete for mates. We found some support for this in our gene expression 323 data. Specifically, we found differential expression in the rac2 and ube3a genes between scale-324 eaters vs generalists. The rac2 gene is associated with the visualization of visible light, 325 metabolism, and behavior (Elsaesser et al. 2010; Goergen et al. 2014). Mutations in the rac2 326 gene affect both male aggression and courtship in Drosophila (Goergen et al. 2014). Differential 327 expression of *ube3a* has also been linked to male aggression. Interestingly, the *ube3a* gene is 328 responsible for producing Ubiquitin-protein ligase E3A, an enzyme that aids in the degradation 329 of proteins, which may be adaptive for the protein-rich diet of scale-eaters which exhibit 330 substantial differential expression of metabolism-related genes (McGirr and Martin 2018). 331 Differential expression of *ube3a* has also been linked to variation in levels of aggression in male 332 rats (Kurian et al. 2007; Stoppel 2014). However, snail-eater and generalist pupfish also adhere 333 to a lekking mating system, although there may be quantitative differences in male competition 334 and degree of lekking among species and lake populations (CHM pers. obs.). 335 The increased female aggression of snail-eaters may also be explained by mating system. 336 Although snail-eaters have been observed mating in the lekking system, not much is known 337 about how their courting behaviors differ from generalists. It is possible that increased levels of 338 female aggression are part of the species' courting ritual. Alternatively, female aggression may 339 have increased incidentally due to selection for decreased maternal care. Our gene expression 340 data indicates that snail-eaters show increased levels of expression for the *nr2e1* gene compared 341 to generalists. This gene (nuclear receptor subfamily 2, group E, member 1) produces a receptor 342 which has been linked to abnormal brain and eye development, as well as increased aggression 343 and lack of maternal care (Young et al. 2002; Abrahams et al. 2005). It is possible that selection 344 for differing levels of maternal care in snail-eaters, compared to generalists or scale-eaters, also

incidentally increased levels of aggression for females. For example, one closely related
outgroup to *Cyprinodon, Jordanella floridae*, exhibits paternal care of eggs through pectoral

347 fanning (St Mary et al. 2004).

348

349 Increased aggression due to indirect selection

350 Alternatively, aggression may have increased via selection on other traits. For example, melanin 351 production and aggression are physiologically linked via the melano-cortin system (Cone 2005; 352 Price et al. 2008). This association has been documented across a wide array of vertebrates and 353 suggests that selection for increased melanin pigmentation in other contexts (e.g. mate choice or 354 camouflage) may incidentally increase aggression (Mcgraw et al. 2003; Ducrest et al. 2008; 355 Price et al. 2008). Indeed, territorial male scale-eating pupfish exhibit jet black breeding 356 coloration, unique among cyprinodontiform fishes, and territorial snail-eating pupfish exhibit the 357 lightest breeding coloration of any Cyprinodon species (Martin and Wainwright 2013a). 358 Similarly, selection for morphological traits may also indirectly increase aggression. We found 359 differential gene expression between scale-eater vs generalist pupfish in the gnaq gene, which is 360 annotated for maternal care (Table 2B). Gnaq is one of four Gq class α -subunits and aids in 361 phospholipase C- β – receptor coupling (Offermanns et al. 1998). Silencing this gene produces 362 severe craniofacial defects in mice, especially in the mandible (Offermanns et al. 1998). C. 363 *desquamator* show extreme craniofacial features, including enlarged oral jaws that may be 364 beneficial for scale-eating. Thus, it is intriguing that selection for increased jaw size may have 365 indirectly selected for increased aggression in this species. Given the enlarged oral jaws of most 366 scale-eating species, this may be a general mechanism indirectly contributing to increased 367 aggression in scale-eaters depending on how frequently this genetic pathway is modified.

368	In conclusion, our data suggest that the aggression hypothesis is not a sufficient					
369	explanation for the origin of scale-eating in pupfish. Instead, increased aggression in both					
370	specialists indicates that aggression may function in dietary specialization or occupation of a					
371	novel niche. Alternatively, increased aggression may be an incidental effect of selection on other					
372	ecological or sexual traits. Specifically, the aggression-boldness syndrome, the melanocortin					
373	system, selection for increased oral jaw size, or metabolic adaptations for increased intake of					
374	protein could all have indirectly increased aggression. Future studies should investigate whether					
375	aggression is adaptive for scale- and snail-eating in pupfish.					
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384	provided logistical support.					
385						
386	Author contributions					
387	MES and CHM conceptualized the project, MES and JAM collected data and performed					
388	analyses, MES wrote the manuscript, and all authors revised the manuscript.					
389						
390	Data accessibility					

- 391 All behavioral datasets from this study will be deposited in the Dryad Digital Repository.
- 392 Transcriptomic raw sequence reads are deposited at the NCBI BioProject database (Title:
- 393 Craniofacial divergence in Caribbean Pupfishes. Accession: PRJNA391309).
- 394

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sexual signal in a parasitized population of field crickets. Biol. Lett. 2:521–524. The Royal
Society.

- 597 **Table 1.** Results of likelihood ratio test for: a) the latency to approach mirror image (mixed-
- 598 effect Cox proportional hazards model); b) the latency to attack mirror image (mixed-effect Cox
- 599 proportional hazards model); and c) the total number of attacks (generalized linear mixed
- 600 model). Significant results are indicated in bold.

model	response variable	predictor	df	χ^2	Р
<i>a</i>) proportional					
hazards	Latency to Attack Mirror Image	Species	3	9.78	0.02
		Sex	1	0.3	0.58
		Species:Sex	3	7.4	0.06
		Species	3	11.21	0.01
b) proportional	Latency to Attack Mirror Image				
hazards		Sex	1	4.29	0.04
		log SL	1	4.8	0.03
		Species:Sex	3	11.12	0.01
	Total Number of	Species	3	7.08	0.07
c) GLMM	Attacks	Sex	1	7.58	0.01
	Attacks	Species:Sex	3	20.91	0.0001
Ļ					

607

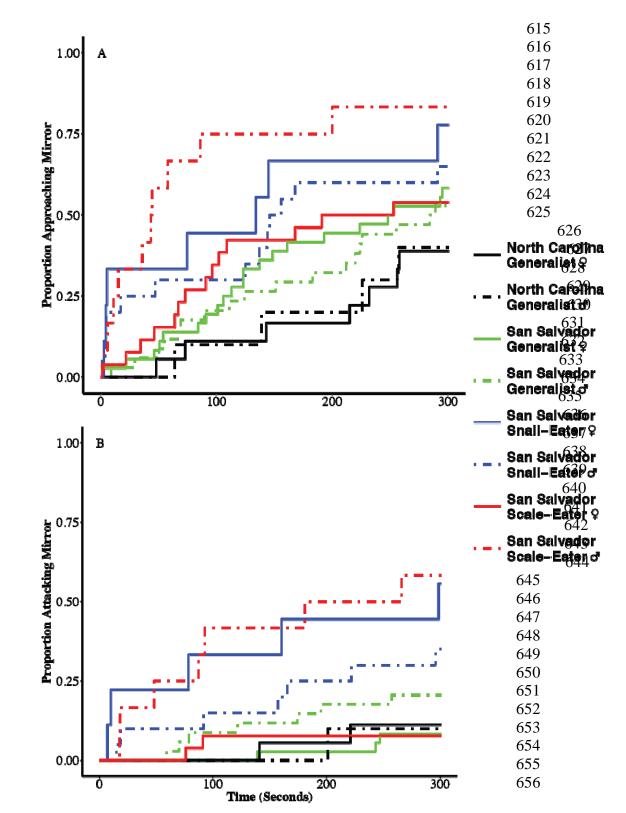
608 **Table 2.** List of all differentially expressed genes in aggression-related pathways between *a*)

- 609 sail-eaters vs generalists and b) scale-eaters vs generalists. 3-6 individuals of each species
- 610 sampled at 8-10 dpf.
- 611

species comparison	gene	log fold change	GO pathway
a) Snail-Eater vs Generalist			
	nr2e1	1.44	Aggression
	heyl	-0.83	Androgen
	rhoa	-0.2	Androgen
	smarca4	0.53	Androgen
	crebrf	-0.7	Maternal Care
b) Scale-Eater vs Generalist			
			Aggression/Maternal Care/
	crhbp	-1.14	Maternal Aggression
			Aggression/Inter-Male
	fkbp14	1.24	Aggression
	penk	1.12	Aggression
	rac2	-0.9	Aggression
	arid1a	0.46	Androgen
	ddx5	0.5	Androgen
	fkbp4	0.62	Androgen
	foxp1	-0.91	Androgen
	heyl	-1.03	Androgen
	igfl	1.22	Androgen
	med12	-0.98	Androgen
	med14	0.74	Androgen
	med16	1.24	Androgen
	rhoa	-0.46	Androgen
	smarca4	0.72	Androgen
	taf1	0.4	Androgen
	ube3a	-0.96	Androgen
	crebrf	-1.41	Maternal Care

dbh	2.02	Maternal Care
gnaq	-1.53	Maternal Care
cul5	0.58	Vasopressin

Figure 1. Time-to-event curves for *a*) the latency to approach mirror image (Cox proportional



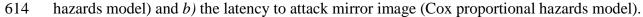


Figure 2. Box plots illustrating the total number of attacks performed by each species and sex (*n*

658 = 165 total individuals tested). Squares represent the total number of attacks for individual

