1 **A behavioral syndrome linking boldness and flexibility facilitates invasion success in**
2 **sticklebacks**
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4 **Running title**: Boldness, flexibility & biological invasions
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17
18 **Keywords**: plasticity, persistence, novel object, intraspecific variation, marine, freshwater,
19 convergent, rapid adaptation
20
21 **Type of article**: Letter
22
23 **Number of words in the abstract** = 149; number of words in the main text (excluding abstract,
24 acknowledgements, references, table and figure legends) = 4427
25
26 **Number of references**: 50
27
28 **Number of figures**: 4
29
30 **Statement of authorship**: All authors agree to submission of the manuscript. MB designed the
31 study, collected and analyzed the data and co-wrote the manuscript. AMB designed the study,
32 contributed to interpretation of the results and co-wrote the manuscript. The authors do not have
33 any conflicts of interest.
34
35 **Data accessibility statement**: Should the manuscript be accepted, the data supporting the results
36 will be archived in Dryad and the data DOI will be included at the end of the article.
37
38

ABSTRACT

For a species to expand its range, it needs to be good at dispersing and also capable of

exploiting resources and adapting to different environments. Therefore, behavioral and cognitive

traits could play key roles in facilitating invasion success. Here, we show that dispersing

sticklebacks are bold, while sticklebacks that have recently established in a new region are

flexible. Moreover, boldness and flexibility are negatively correlated with one another at the

individual, family and population levels. Multiple lines of evidence suggest that the divergence

in boldness and flexibility is likely to be evolutionary in origin. If boldness is favored in invaders

during the initial dispersal stage, while flexibility is favored in recent immigrants during the

establishment stage, then the link between boldness and flexibility could generate positive

correlations between successes during both dispersal and establishment, and therefore play a key

role in facilitating colonization success in sticklebacks and other organisms.

INTRODUCTION

Understanding the factors that allow a species to expand its range and adapt to changing habitats is increasingly important in the face of anthropogenic change. Natural biological invasions can reveal how and why certain organisms can excel in response to novel selection pressures (Whitney & Gabler 2008). In addition to the importance of propagule pressure (Lockwood *et al.* 2005), stochasticity, and the opening of new niches at the edge of species boundaries, there is growing evidence that particular traits (e.g., r-selected life histories (Capellini *et al.* 2015), habitat breadth (Blackburn *et al.* 2009), large brains/cognitive abilities (Sol *et al.* 2005)) might promote biological invasions.

Behavioral and cognitive traits are likely to play an important role in allowing animals to move into and become established in new environments (Sih *et al.* 2010; Wright *et al.* 2010; Hoset *et al.* 2011; Chapple *et al.* 2012). For example, behavioral plasticity allows animals to rapidly adjust their phenotype in response to changes in environmental cues (Wolf *et al.* 2008; Coppens *et al.* 2010; Sih *et al.* 2012). Cognitive processes – how animals perceive, process and retain information about their environment and then use that information to make decisions (Shettleworth 2010) – may play an especially important role during biological invasions because they influence the ability of animals to enter new habitats, exploit new niches, become established and spread (Wright *et al.* 2010; Griffin *et al.* 2016; Szabo *et al.* 2020). For example, recent immigrants have to be willing to approach and interact with novel stimuli in order to gain information about their new environment (i.e., neophilia (Mettke-Hofmann *et al.* 2009)). Additionally, previously successful behavioral patterns may no longer be successful in new environments, so immigrants need to be able to stop persisting on ineffective responses and flexible enough to attempt new approaches (Griffin *et al.* 2016).

Successfully colonizing a new environment can be broken down into discrete stages, e.g.,

dispersal, colonization, establishment and spread (Chapple *et al.* 2012), and invasion success

likely relies on different behavioral and cognitive traits in each stage (Sih *et al.* 2012). For

example, dispersers need to be bold, while immigrants can succeed in a new environment when

they are willing to investigate novel stimuli and when they are able to quickly inhibit old and

ineffective behaviors, i.e., if they are flexible (Sih *et al.* 2012). If boldness, neophilia and

behavioral flexibility are no longer beneficial or even costly once a population becomes

established in a new environment, they may be lost (Wright *et al.* 2010; Herczeg *et al.* 2020).

Numerous studies have documented differences in behavioral traits between invading and

established populations, e.g., (Martin & Fitzgerald 2005; Pintor *et al.* 2008; Damas-Moreira *et*

al. 2019; Cohen *et al.* 2020), and a handful of studies have shown that they have a heritable

basis, e.g., (Sargent & Lodge 2014; Gruber *et al.* 2017). If there is an underlying genetic basis to

the behavioral and cognitive traits that facilitate range expansions or biological invasions, then

we might expect those traits to evolve over the course of an invasion. According to this

hypothesis, behavioral and cognitive traits important for invasion should vary in a systematic

way between dispersing populations compared to newly-derived and well-established

populations when they are reared in a common garden. Moreover, if invasion success requires

different behavioral and cognitive traits in the different invasion stages, then mechanisms that

package these traits together could be key to the success of invasive species (Sih *et al.* 2012). For

example, traits important for the dispersal and establishment stages are coupled together in

western bluebirds, and this facilitates the expansion of their range (Duckworth & Badyaev 2007).

Threespined sticklebacks (*Gasterosteus aculeatus*) are a model system for studying trait

evolution during biological invasions. Throughout their evolutionary history, marine sticklebacks

have repeatedly colonized freshwater environments, rapidly adapted to them and diversified

(Stuart *et al.*). Sticklebacks can also spread and have dramatic impacts on freshwater

communities (Eklöf *et al.* 2020; Gugele *et al.* 2020). Work on this system has primarily focused

on a suite of morphological and physiological traits that repeatedly evolves when marine

sticklebacks invade freshwater habitats (Bell *et al.* 2004; Colosimo *et al.* 2005), with evidence

that haplotypes containing a set of coadapted alleles are maintained at low frequency in the

ocean and are repeatedly tapped during adaptation to freshwater (Bassham *et al.* 2018).

However, little is known about the behavioral and cognitive mechanisms that may facilitate the

invasion of sticklebacks into new habitats and which may evolve once a population becomes

established (Foster *et al.* 2015).

We took advantage of the repeated invasion of freshwater habitats by sticklebacks by

comparing marine stickleback to sticklebacks from recently-derived and well-established

freshwater populations in Alaska, with replicate populations of each type. For each population,

we reared clutches of field-fertilized eggs in a controlled laboratory environment and scored

individuals for three informative and repeatable behavioral traits: boldness, behavioral flexibility,

and neophilia (Bensky & Bell 2020). We hypothesized that boldness is favored in dispersers, that

neophilia and behavioral flexibility are favored in newly-arrived immigrants, and that these traits

may be subject to relaxed selection and possibly lost once a population becomes established. The

replicate populations allow us to assess the generality of the patterns and comparing the newly-

established to well-established populations provides insight into whether different traits are

favored during initial establishment versus population persistence (Merwin 2019). Rearing

animals under common-garden conditions in the lab also allowed us to determine if behavioral

variation within and among populations likely reflect evolved, genetically based differences.

MATERIALS AND METHODS

Adult sticklebacks were collected via minnow traps from six populations ranging from

the Matanuska-Susitna Valley to the Kenai Peninsula of Alaska (Figure S1; Table S1) during

June 2017. Two populations (Rabbit Slough and Resurrection Bay) occur in the ancestral marine

environment, while the remaining four populations occur in freshwater. Two of the freshwater

populations (Big Beaver and Cornelius, hereafter referred to as "well-established") are derived

from natural colonization events that presumably occurred hundreds to thousands of years ago,

after the last glacial maximum, while the other two freshwater populations are recently derived

via natural recolonization (Loberg: 28-34 years prior to collection; (Bell *et al.* 2004)) or

experimental seeding (Cheney: 8 years prior to collection; (Bell 2016)). As is typical for

sticklebacks, the marine populations are only weakly genetically differentiated from each other

($F_{ST} = 0.0076$; (Hohenlohe *et al.* 2010)) while the freshwater populations are more strongly

genetically differentiated (M. Bell, K Veeramah, personal communication).

Eggs were fertilized in the field following previously established protocols (see (Wund *et*

al. 2012)). Two to three days post fertilization, the eggs were transferred to 50 mL canonical

tubes and shipped overnight in coolers filled with ice packs to the University of Illinois Urbana-

Champaign where they were raised in common garden conditions in the lab. Artificial

incubation controls for environmental paternal effects due to receiving paternal care therefore it

is likely that phenotypic differences among the lab-reared populations reflect heritable

differences (although environmental maternal effects could also contribute).

Clutches were reared in separate tanks (9.5L 32 x 21 x 19 cm) where the embryos were

incubated in a cup with a mesh bottom and placed over an air bubbler. Fish were kept at 60°F

with an even light cycle (12L:12D) for the entirety of the experiment. All families were kept on

one of two recirculating flow-through water racks, which consisted of a series of particular,

biological, and IV filters and had three different shelves (Aquaneering , San Diego, USA). 10%

of each tank's water was replaced each day. Family tank position was pseudo-randomly assigned

so that all populations were evenly distributed across both racks and the three levels of shelves.

Importantly, we elected to rear the fish and measure their behavior in freshwater (~5ppt), thereby

simulating the conditions that marine sticklebacks encounter when they move into freshwater.

Because the marine populations studied here are naturally anadromous (Rabbit Slough: (Bell

2016); Resurrection Bay: R. King, pers. comm.), i.e., they spawn in fresh/brackish waters, their

early offspring development typically occurs under low salinity.

Behavioral assays

Summary of the behavioral assays. Neophilia was measured as response to a novel

object. Boldness was measured as latency to emerge from a refuge, a reliable and widely-used

behavioral assay in fishes (Wilson & Godin 2009; Pearish *et al.* 2013). Boldness was quantified

at the individual level as average latency to emerge across four independent trials. Behavioral

flexibility was measured in a barrier task: after pretraining individuals to expect a food reward

upon emergence from a refuge, individuals were confronted by a transparent barrier that they had

to swim around in order to get the food reward. Individuals that continue to follow the prepotent

search pattern established during training spend relatively more time at the point of the barrier

closest to the food reward ("barrier apex"), which we interpret as relatively low flexibility. In

contrast, individuals that quickly break away from the previously-established behavior pattern

spend relatively little time spent at the apex of the barrier, which we interpret as relatively high

flexibility. A previous study in sticklebacks found that time at the apex of the barrier predicts

reversal learning performance (Bensky & Bell 2020), another common metric of behavioral

flexibility.

Detailed experimental methods. 18 observation tanks (36L x 33W x 24H cm) were used

for behavioral testing. When the fish were approximately eight months of age (approximately 40

mm standard length), the testing phase of the experiment began. Behavioral assays were carried

out over the course of 5 months. Families, sexes and populations were measured in a

pseudorandomized order, such that male and female offspring from different families and from

different populations were measured in the same block. Individuals were randomly selected from

each family, and their weight and length were measured. During the testing phase of the

experiment, fish were only fed during the behavioral tests to help maintain motivation.

Acclimation phase

In order to ensure that an individual had acclimated to their home tank and was motivated

to eat during the behavioral tests, the individual was presented with food via a petri dish at the

center of their home tank, and the individual had to eat the food within 10 minutes on three

consecutive days in order to proceed to the next step. On average, it took 5.1 days for fish to

meet this criterion (range = 3 to 17 days).

Novel object test (neophilia)

Individuals' response to a novel object (toy lion; 10L x 7H cm; TERRA by Battat,

Montreal, Canada) was recorded the day after the fish met criterion in the acclimation phase. The

toy lion was selected as a novel object because the fish had no prior experience with this object,

there was no presumed evolutionary history with the object's shape, and it was made up of

neutral colors.

The individual's behavior was measured for five minutes after their first approach to the

novel object (i.e., first time within one body length of the novel object and oriented directly

towards it). We interpret more time spent near and oriented towards the novel object as greater

time investigating the object (i.e., higher neophilia).

Latency to emerge (boldness)

For each individual, we recorded their latency to emerge from a refuge on their first four

training trials for the barrier task (described below) and used the average of those measures as a

proxy for boldness (Movie S1). Latency to emerge was repeatable across the four trials within

each population (r=0.56-0.72, Table S4).

Barrier detour task (flexibility)

Pretraining for the barrier detour task started on the same day the novel object test was

completed. The goals of pretraining were to train the individual to learn that there would

consistently be a food reward in the middle of the tank, establish food motivation in this context,

and create a prepotent response of leaving a shelter to directly approach and eat the food reward.

During pretraining, individuals were trained for one session per day, with each session

comprising four trials. To start the trial, the observer removed a cork from the side of the shelter

and the fish was given ten minutes to exit. Upon exiting the fish was allowed five minutes to eat

the worm.

After eating the worm, the fish was placed back into the shelter in preparation for the

next trial. If the fish did not emerge from the shelter within ten minutes after the cork was pulled

or eat within five minutes after emergence, the observer recorded the maximum times for these

behaviors, removed the food reward and gently poured the fish out of the shelter if necessary.

Latency to emerge from a refuge during the first pretraining session was used as a measure of

boldness (see above).

Training for the barrier task was criterion based. In order to move on to the barrier task

following pretraining, the individual had to emerge from the shelter within 10 minutes and

directly approach and eat the food reward within five seconds on three out of the four trials. The

one failed attempt could not be on the fourth trial; this requirement ensured that the fish would

be motivated throughout the four trials. Fish were given a maximum of four days to reach

criterion. Fifteen fish did not meet criterion (Big Beaver: 7, Cornelius: 1, Loberg: 1, Cheney: 3,

Rabbit Slough: 2, Resurrection Bay: 1).

Once an individual met criterion, the individual moved on to the barrier detour task the

following day. This task also comprised four trials. The first two trials were exactly the same as

the pretraining trials in order to reinforce the direct search pattern. On the third trial a transparent

semi-circular barrier was placed between the shelter and food reward. The opening into the

barrier area was positioned directly in front of the entrance to the shelter. After removing the

cork the individual was allowed 30 minutes to emerge from the shelter, navigate around the

barrier and eat the food reward (Movie S2). The observer recorded the duration of the first bout

(no break in contract longer than five seconds) at the apex of the barrier. In order to confirm that

the fish that spent little time at the barrier apex during the third trial were still motivated to eat,

the fish's behavior was observed for a fourth trial during which no barrier was present.

Altogether, a total of n=262 individuals from n=8-11 families per population (n=2-7 full

sibs per family) completed the novel object test and boldness assay. A total of n=247 individuals

from n=8-11 families per population (n=1-4 full sibs per family) completed the barrier task. The

experiments were approved by the Institutional Animal Care and Use Committee (IACUC) of

the University of Illinois Urbana-Champaign (IACUC protocol #15077).

Statistical analysis

R 3.5.3 (http://www.r-project.org/) was used for statistical analyses. Positively-skewed

variables were log-transformed to improve normality, model residuals were also visually

inspected for deviations from normality.

We used linear mixed models (Team 2013)(package = "lme4"; function = "lmer") to

examine the behavioral data in each assay separately. We created models in which population

and sex were included as fixed factors, and body length was included as a covariate. FamilyID

nested within population was included as a random variable. The statistical significance of the

effect of FamilyID was assessed by AIC (Akaike 1973), i.e., by comparing models with and

without the effect of FamilyID (R Core Team 2016; package = "care"; function = "anova"). We

infer that a trait has an underlying heritable basis when the trait differs among populations

(because they were reared in a common garden), and/or when FamilyID improves model fit. To

further explore the heritable basis to these traits we also computed broad-sense heritabilities and

genetic correlations within each population following (Hadfield 2010; Wilson *et al.* 2010). We

used weakly informative inverse-gamma priors for the 'residual' and 'genetic' effects (by setting

the MCMCglmm parameters $V=1$, $nu=0.002$). In order to test for genetic correlations, a

2x2 covariance matrix was specified with the degree of belief parameter set to n=1.002. The raw

positively-skewed duration data for all the behavioral measures were used for this analysis, and

thus all measures were rounded to the nearest integer and a Poisson distribution was used. The

posterior distribution was sampled every 3000 times (thinning interval), following a burn-in

period of 100□000 iterations with a total run of 10□000□000 iterations. The relatively modest

number of families and full sibs within each family resulted in large confidence intervals around

the estimates therefore we interpret those results with some caution.

To examine the structure of the phenotypic variation both among and within populations,

we examined correlations between traits at the individual, family and population level.

See SI Methods for more details.

RESULTS

Sticklebacks from dispersing populations were bold

There was significant variation among populations in boldness (latency to emerge from a

refuge ($F_{5,59}$ = 3.30, p = 0.01, Table S1, Figure 1)). Sticklebacks from dispersing populations

(marine) were more bold than fish from well-established freshwater populations (z = 2.745, n =

262, p = 0.006). The two recently-derived populations provide insight into the rate at which

boldness diverged from the ancestral marine behavioral type. Sticklebacks from Cheney Lake, a

population that was established eight years prior to collection, resembled marine populations and

differed from the well-established freshwater populations (Cheney vs. well-established: z = -

2.497, n = 262, p = 0.034). In contrast, sticklebacks from Loberg Lake, a population that was

established 28-34 years prior to collection, more closely resembled the well-established

freshwater populations (Loberg Lake vs marine: z = 2.429, n = 262, p = 0.040). After correcting

for multiple comparisons, there was a trend for sticklebacks from Cheney Lake to emerge faster

compared to sticklebacks from Loberg Lake (z = -2.249, n = 91, p = 0.064).

There was significant variation in boldness among families within populations (FamilyID

improved model fit (X^2 = 9.394, df = 1, p = 0.002; AIC_{with} = 774.50; $AIC_{without}$ = 781.89)).

Estimates of broad sense heritability of boldness in the six populations ranged from 0.1 to 0.75

(average = 0.48) and was significantly different from zero in the Big Beaver, Cornelius and

Resurrection Bay populations (Table S2).

Sticklebacks from established populations were flexible

There was significant variation among populations in behavioral flexibility (time at the

apex of a barrier ($F_{5,58}$ = 3.495, p = 0.008, Table S1, Figure 2)). Sticklebacks from the well-

established populations were more flexible (spent less time at the apex of a barrier), compared to

sticklebacks from the dispersing marine populations (z = -2.175, n = 158, p = 0.030). As was the

case for boldness, the two recently-derived populations show different patterns: flexibility in

Cheney Lake – the most recently established freshwater population – resembled flexibility in the

dispersing populations, while flexibility in Loberg Lake – the freshwater population that was

established 28-34 years prior to collection – resembled flexibility in the well-established

freshwater populations (Cheney Lake vs Loberg Lake: z = 2.821, n = 87, p = 0.013, Cheney

Lake vs well-established: z = 3.373, n = 131, p = 0.002).

There was significant variation among families within populations in flexibility (X^2 =

7.350, df = 1, p = 0.007; AIC_{with} = 723.74; $AIC_{without}$ = 729.09). Estimates of broad sense

heritability of flexibility in the six populations ranged from 0.22 to 0.86 (average = 0.45) and

was significantly different from zero in the Cornelius population (Table S2).

Neophilia did not evolve systematically during the invasion

Neophilia (time near and oriented to a novel object) did not vary among populations or

between the sexes (Table S1) and did not vary in a systematic manner among the different types

of populations (Figure 3). Larger fish were less neophilic (β = -0.033, t = -2.631, df = 247.79,

p=0.009, Figure S2, Table S1). We did not detect variation among families within populations in

neophilia, as FamilyID did not significantly improve model fit (X^2 = 2.369, df = 1, p = 0.124;

AIC_{with} = 636.57; $AIC_{without}$ = 636.94). Estimates of broad sense heritability of neophilia in the

six populations ranged from 0.09 to 0.87 (average = 0.34) and was significantly different from

zero in the Cornelius population (Table S2).

A boldness-flexibility syndrome facilitates invasion success

Given that sticklebacks from the dispersing populations were bolder and less flexible

compared to sticklebacks from established populations, we tested how boldness and flexibility

were correlated with one another. Consistent with the pattern described above, individuals that

were more bold (quickly emerged from the refuge) were less flexible (spent more time at the

barrier apex; r = -0.522, n = 245, p<0.001, Table S3), and this pattern was also evident at the

family (r = -0.688, n = 62, p<0.001) and population (r = -0.91, n = 6, p=0.01) levels (Figure 4).

The average genetic correlation between boldness and flexibility within the six populations was r

= -0.69 (range: -0.59 to -0.77) and was significantly different from zero in the Cornelius

population (Table S3).

DISCUSSION

The possibility that behavioral and cognitive traits might facilitate and evolve during

natural biological invasions is intriguing but difficult to study directly. Here, we took advantage

of a model system for invasions to test the hypothesis that boldness is favored in dispersers, that

neophilia and flexibility are favored in recently-arrived immigrants and that these traits are

subject to relaxed selection and possibly lost once a population becomes established in a new

environment. We found that boldness and flexibility evolve in a systematic way when marine

sticklebacks colonize freshwater habitats. Specifically, the dispersing populations were bold, and

well-established populations were flexible. These traits varied between the two recently

established populations relative to time since establishment. Moreover, boldness and flexibility

were negatively correlated with one another at the individual, family and population levels.

Differences in boldness and flexibility were evident in a common garden environment, there was

significant variation among families in both of these traits, and estimates of their heritability and

the genetic correlation between them were relatively high for behavioral traits (Dochtermann *et*

al. 2019). These lines of evidence suggest that there is a heritable component to the traits and

that their divergence is likely to be evolutionary in origin.

If boldness is favored in invaders during the initial dispersal stage, while flexibility is

favored in recent immigrants during the establishment stage, then a tight link between boldness

and flexibility could generate positive correlations between successes during both the dispersal

and establishment stage, and therefore play a key role in facilitating invasion success in this

species (Sih *et al.* 2012). Selection favoring boldness in dispersers and flexibility in immigrants

could cause the population to move along a ridge of high fitness whenever sticklebacks diverge

from the ancestral marine behavioral type (Figure 4). An outstanding question is what maintains

the extensive phenotypic and genetic variation in boldness and flexibility within the populations

(Figures 1-4). Selection might not be strong enough to effectively purge the variation, and/or

there may be ongoing gene flow between marine and freshwater habitats which maintains

recessive alleles associated with behavioral flexibility at low frequency in marine populations

(transporter hypothesis (Schluter & Conte 2009)), as appears to operate at the *Eda* locus (Jones *et*

al. 2012) and surrounding genomic regions (Bassham *et al.* 2018).

We found no support for the hypothesis that traits which promote dispersal and early

establishment in a new environment are lost once a population becomes well-established (Wright

et al. 2010), i.e., no evidence that newly-derived populations were more bold, neophilic and

flexible than well-established populations. One possible explanation for this pattern is that

assortative mating within dispersing populations causes dispersing behavioral phenotypes to be

maintained early in the establishment process, even in the presence of selection favoring more

flexible phenotypes (Shine *et al.* 2011). The comparison between the two newly-derived

populations (Cheney Lake and Loberg Lake, respectively) is also insightful; the two newly

derived populations either tended to resemble the dispersing populations or the well-established

populations. Cheney Lake was founded more recently than Loberg Lake (9 versus 28-34 years

prior to this study); if it takes longer than 10 generations for behavioral and cognitive traits to

diverge from the ancestral marine behavioral type then *time since establishment* could be

important. Alternatively, or in addition, the phenotypic differences between sticklebacks from

Loberg Lake and Cheney Lake could reflect differences in the way that the two lakes were

colonized; Loberg Lake was naturally colonized, while sticklebacks were experimentally

introduced to Cheney Lake. If particularly flexible individuals were more likely to disperse into

Loberg Lake, but a random sample of behavioral types were artificially introduced into Cheney

Lake (Cote *et al.* 2010; Edelaar & Bolnick 2012; Canestrelli *et al.* 2016), then *non-random*

dispersal could be contributing to the rapid evolution of increased flexibility in Loberg Lake.

Given evidence from the literature (Cote *et al.* 2010; Canestrelli *et al.* 2016) and from this study

that bold individuals are more likely to disperse, and that boldness and flexibility are tightly

negatively correlated with one another, this explanation seems unlikely. Further studies tracking

how behavioral and cognitive traits change over time in the Cheney Lake population (and similar

experimental lakes (e.g., Scout Lake)) could help discriminate between the assortative mating,

time since establishment and non-random dispersal hypotheses.

We originally hypothesized that neophilia would be favored in recently-derived

populations because seeking and/or being willing to investigate novel stimuli may help newly-

arrived immigrants locate new habitats and discover novel resources, but we found no support

for this hypothesis. One possible explanation for the failure to find systematic differences in

neophilia among the populations is that neophilia may actually be disadvantageous in a new

environment because it can expose animals to dangerous stimuli they have never encountered

before. Another potential (nonexclusive) explanation based on our results is that neophilia may

not evolve as readily because it may be less heritable (effect of FamilyID was nonsignificant,

lower H^2 estimate). Instead, neophilia may be more influenced by age or experience: in this

study, smaller (and younger: $r = -0.159$, $t = -2.595$, $n = 262$, $p=0.01$) fish were more neophilic,

which is consistent with other studies which have shown that novelty-seeking decreases with age

(Stansfield & Kirstein 2006).

Sticklebacks are a powerful model system for understanding how and why certain traits

repeatedly evolve whenever organisms invade new habitats. Accumulating evidence suggests

that sticklebacks have evolved mechanisms for rapidly adapting to new environments with

alleles conferring the freshwater-adapted phenotype maintained at low frequency in the ocean

(Colosimo *et al.* 2005). But the role of behavior and cognition in facilitating evolutionary

processes in this system has received less attention (Foster *et al.* 2015). Our results suggest that a

behavioral mechanism – a behavioral syndrome linking boldness and flexibility together – can

contribute to rapid adaptation in this and other organisms.

ACKNOWLEDGEMENTS

We thank Matt Wund, Mike Bell, John Baker, and the members of the Foster Lab for

help with field work. The Bell lab, Andy Suarez, Jason Keagy, Becky Fuller and Katy Heath

provided valuable feedback. Financial support was provided by the National Science

Foundation's Integrative Graduate Education and Research Traineeship program and by the

University of Illinois. This material is based upon work supported by the National Science

Foundation under Grant No. IOS 1121980 and by the National Institutes of Health under award

number 2R01GM082937-06A1.

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540

542 **FIGURES**

544 **Figure 1**. Variation among populations in boldness (latency to emerge from a refuge). Smaller

545 values indicate greater boldness. Figure shows boxplots with individual data points

546 superimposed.

Figure 2. Variation among populations in flexibility (time at the apex of the barrier). Smaller

values indicate greater flexibility, i.e., less time persisting on a previously-successful behavior

pattern. Figure shows boxplots with individual data points superimposed.

560 **Figure 3**. Variation among populations in neophilia (time near and oriented to the novel object).

561 Figure shows boxplots with individual data points superimposed.

Figure 4. Relationship between flexibility (time at the barrier apex) and boldness (latency to

emerge) within and among populations. Dispersers are relatively bold (emerge quickly) while

immigrants are relatively flexible (spend less time at the barrier apex). Large circles represent

populations means, medium circles show family means, small circles show individual data

points. Circles are color coded by population. The line and text drawn on the figure are for visual

purposes only.

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