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# Factors affecting individual foraging specialization and temporal diet stability across the range of a large "generalist" apex predator

Adam E. Rosenblatt Department of Biological Sciences, Marine Sciences Program, Florida International University, arosenbl@fu.edu

James C. Nifong *Unviersity of Florida* 

Michael R. Heithaus Department of Biological Sciences and Marine Sciences Program, Florida International University, heithaus@fiu.edu

Frank J. Mazzotti University of Florida

Michael S. Cherkiss U.S. Geological Survey

See next page for additional authors

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#### Authors

Adam E. Rosenblatt, James C. Nifong, Michael R. Heithaus, Frank J. Mazzotti, Michael S. Cherkiss, Brian M. Jeffery, Ruth M. Elsey, Rachel A. Decker, Brian R. Silliman, Louis J. Guillette Jr., Russell H. Lowers, and Justin C. Larson

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4	Adam E. Rosenblatt <sup>1,*</sup> , James C. Nifong <sup>2</sup> , Michael R. Heithaus <sup>1,3</sup> , Frank J. Mazzotti <sup>4</sup> , Michael S.
5	Cherkiss <sup>5</sup> , Brian M. Jeffery <sup>4</sup> , Ruth M. Elsey <sup>6</sup> , Rachel A. Decker <sup>1</sup> , Brian R. Silliman <sup>2</sup> , Louis J.
6	Guillette Jr. <sup>7</sup> , Russell H. Lowers <sup>8</sup> , Justin C. Larson <sup>2</sup>
7	
8	Affiliations
9	<sup>1</sup> Department of Biological Sciences, Marine Sciences Program, Florida International University,
10	North Miami, FL 33181, USA
11	<sup>2</sup> University of Florida, Department of Biology, Gainesville, FL 32611, USA
12	<sup>3</sup> School of Environment, Arts, and Society, Florida International University, North Miami, FL
13	33181, USA
14	<sup>4</sup> Fort Lauderdale Research and Education Center, University of Florida, Davie, FL 33314, USA
15	<sup>5</sup> U.S. Geological Survey, Southeast Ecological Science Center, Davie, FL 33314, USA
16	<sup>6</sup> Louisiana Department of Wildlife and Fisheries, Rockefeller Wildlife Refuge, Grand Chenier,
17	LA 70643, USA
18	<sup>7</sup> Medical University of South Carolina, Department of Obstetrics and Gynecology and Hollings
19	Marine Laboratory, Charleston, SC 29412, USA
20	<sup>8</sup> InoMedic Health Applications, Kennedy Space Center, FL 32899, USA
21	*Corresponding author: email: arose007@fiu.edu, phone: 215-913-4078, fax: 305-919-4030 <sup>1</sup>

<sup>&</sup>lt;sup>1</sup> Author Contributions: AER, JCN, MRH, and FJM conceived the idea for the study. AER, JCN, MSC, BMJ, RME, RAD, LJG, RHL, and JCL collected the data. AER and RAD analyzed the data. AER, JCN, MRH, and BRS wrote the manuscript; all other authors provided editorial advice.

22 Abstract

23 Individual niche specialization (INS) is increasingly recognized as an important component of ecological and evolutionary dynamics. However, most studies that have 24 investigated INS have focused on the effects of niche width and inter- and intraspecific 25 competition on INS in small-bodied species for short time periods, with less attention paid to 26 INS in large-bodied reptilian predators and the effects of available prey types on INS. We 27 investigated the prevalence, causes, and consequences of INS in foraging behaviors across 28 different populations of American alligators (Alligator mississippiensis), the dominant aquatic 29 30 apex predator across the southeast US, using stomach contents and stable isotopes. Gut contents revealed that over the short-term, although alligator populations occupied wide ranges of the INS 31 spectrum, general patterns were apparent. Alligator populations inhabiting lakes exhibited lower 32 INS than coastal populations, likely driven by variation in habitat type and available prey types. 33 Stable isotopes revealed that over longer time spans alligators exhibited remarkably consistent 34 use of variable mixtures of carbon pools (e.g., marine and freshwater food webs). We conclude 35 that INS in large-bodied reptilian predator populations is likely affected by variation in available 36 prey types and habitat heterogeneity, and that INS should be incorporated into management 37 38 strategies to efficiently meet intended goals. Also, ecological models, which typically do not consider behavioral variability, should include INS to increase model realism and applicability. 39 40

41 <u>Key words:</u> American alligator, *Alligator mississippiensis*, stomach content analysis, stable
42 isotope analysis, food web

#### 43 Introduction

Intrapopulation foraging specialization can be attributed to differences between sexes 44 ("ecological sexual dimorphism"; Temeles et al. 2000), morphological types ("resource 45 polymorphisms"; Skulason and Smith 1995), and age groups ("ontogenetic niche shifts"; Polis 46 1984). Increasingly, however, it is recognized that individuals within a population can exhibit 47 48 considerable variation in trophic interactions that are not attributed to these factors, but instead to individual niche specialization (INS) that may be caused by differences in learning, 49 morphological and physiological adaptive plasticity, and genetic and epigenetic expression (Dall 50 51 et al. 2012). Individual niche specialization has important potential implications for evolutionary 52 processes (Knudsen et al. 2010) and community and population dynamics (Bolnick et al. 2003; Dall et al. 2012). For example, some sea otter (Enhydra lutris) populations consist of individuals 53 that exhibit extreme dietary specializations, possibly for their entire lives, likely resulting in low 54 intraspecific competition and variable responses of individuals to food web perturbations (Estes 55 et al. 2003). 56

A number of studies have examined various factors that affect the magnitude or 57 occurrence of INS, including niche size (Bolnick et al. 2007; Woo et al. 2008; Araujo et al. 2009; 58 59 Darimont et al. 2009), intraspecific competition (Estes et al. 2003; Tinker et al. 2008), and interspecific competition (Bolnick et al. 2010). However, another variable, prey community 60 composition, has not been thoroughly examined as a potential factor affecting INS. Prey 61 62 community composition is important to consider in the context of INS because for some species consumption of different prey types involves different handling times and attack success rates 63 (Holling 1959; Kislalioglu and Gibson 1976). For example, if a predator population's habitat 64 65 contains abundant easy-capture prey (e.g., gastropods), then individuals in the population could

66 all consume large numbers of that prev group because of short handling times and high attack success rates. Such foraging behavior would cause each individual to exhibit dietary patterns 67 very similar to conspecifics, resulting in low INS. In contrast, a habitat containing few easy-68 capture prey and different types of hard-capture prey (e.g., mammals and birds) could result in 69 higher INS because the predators are less likely to all focus on the same elusive prey group and 70 experience similar attack success rates (e.g., Baird et al. 1992). Killer whales (Orcinus orca) in 71 the eastern North Pacific Ocean may fit such a pattern: two distinct groups of killer whales feed 72 on different prey types, with "transients" primarily consuming pinnipeds and "residents" 73 74 primarily consuming fish (Baird et al. 1992). For killer whales, pinnipeds are more difficult to capture and require more handling time than fish (Baird et al. 1992), and the resulting dietary 75 specialization patterns of the two populations are highly divergent: residents display low dietary 76 variability with prey killed by residents dominated by one genus of fish (Oncorhynchus) which 77 makes up 98% of the diet, while transients display higher dietary variability with prey killed by 78 transients distributed over 10 genera with harbor seals (*Phoca vitulina*) making up 55% of the 79 80 diet (Ford et al. 1998).

The effects of prey community composition on INS are particularly important to 81 82 investigate for large apex predators because they generally roam widely and can inhabit multiple ecosystems with different types of prey communities. Furthermore, large apex predator 83 populations are generally declining globally (Ripple et al. 2014) and understanding INS patterns 84 85 could be important for crafting appropriate conservation strategies (Bolnick et al. 2003). Some studies have documented the presence of INS in apex predators (Baird et al. 1992; Estes et al. 86 87 2003; Tinker et al. 2008; Darimont et al. 2009; Matich et al. 2011; Thiemann et al. 2011), but 88 few have addressed INS in one species across habitats with different prey community

compositions over large spatial scales. If there is variation in the strength of INS among different populations of the same top predator species across space and time (e.g., Baird et al. 1992), then the effects of those top predators on lower trophic levels could vary considerably within and among populations and their roles may differ markedly across ecosystems.

93 Here we investigate the prevalence and stability of INS as well as the effects of prev 94 community composition on INS in a well-studied large apex predator: the American alligator (Alligator mississippiensis). Using complementary techniques, stomach contents analysis (SCA; 95 provides short-term dietary data) and stable isotope analysis (SIA; provides long-term dietary 96 97 data), we assess INS in alligators across their range and a variety of habitats. Alligators are excellent model "generalist" apex predators for such a study because: 1) their diets have been 98 examined across their range repeatedly, 2) they inhabit every type of fresh and brackish water 99 100 habitat across the southeastern US (Mazzotti and Brandt 1994), and 3) their diets at the population level are typically highly diverse. We hypothesized that alligator populations 101 inhabiting lakes would display relatively low degrees of INS because of low habitat variability 102 103 and prey communities containing many easy-capture prey (Darby et al. 2006). Conversely, we hypothesized that alligator populations in coastal habitats that have access to a variety of distinct 104 105 habitat types (e.g., freshwater marshes, dynamic estuarine zones, marine areas) and to fewer 106 easy-capture prey would exhibit higher degrees of INS.

#### 107 Materials and methods

#### 108 Stomach contents collection and analyses

We compiled alligator stomach contents data from seven published studies containing 109 data collected from 1220 alligators between 1977 and 2004 and five new datasets collected from 110 232 alligators between 2007 and 2012 (Table 1). The datasets included samples from a wide 111 112 geographic range and many habitat types (freshwater lakes, mangrove rivers, salt marshes, barrier islands; Fig. 1). In some studies stomachs were sampled as part of state control programs 113 (e.g., state-sanctioned hunting and nuisance alligator removal programs), while in others data 114 115 were collected non-lethally using the hose-Heimlich technique (Table 1; Fitzgerald 1989). In studies that examined the technique's efficacy, 100% of ingested prey items were recovered from 116 91% of the alligators tested (Fitzgerald 1989; Rice et al. 2005; Nifong et al. 2012); therefore, we 117 assumed no sampling bias between studies that used lethal or non-lethal methods. Prey items 118 found in the stomach contents were classified to the lowest possible taxon either immediately 119 after collection or after preservation. 120

121 To assess the prevalence of INS in alligator populations we applied Roughgarden's (1972, 1979) concept of total niche width (TNW; full range of food resources used by a 122 123 population), which is subdivided into a between-individual component (BIC; variance in food resource use between individuals) and within-individual component (WIC; variance in food 124 resource use within individuals), such that TNW = BIC + WIC. If BIC > WIC for a given 125 126 population, then the diets of individuals are more diverse in comparison to each other than they are diverse within each individual. We divided BIC by TNW to generate an index of 127 specialization that varied between 0 and 1, where 0 = pure generalist population (individuals 128 129 completely overlap with population's resource use) and 1 = pure specialist population

(individuals do not overlap at all with other individuals in the population). We chose to focus on
the BIC/TNW index because it is simple to calculate and because other INS metrics generally
produce similar results (Bolnick et al. 2002). Calculations of BIC/TNW are biased by the
inclusion of individuals that only contain prey items from one prey group in their stomach
(Bolnick et al. 2002), so we applied sorting rules to each dataset to limit bias in our INS results
and produce more conservative BIC/TNW values (explanation in electronic supplemental
materials).

137 Individual specialization metrics like BIC/TNW are difficult to compare across 138 populations without accounting for variable numbers of available prey types across different 139 habitats. Thus, to standardize our BIC/TNW values we converted them into adjusted E values 140 ( $E_{adj}$ ; Araujo et al. 2011) using the equation

141 
$$E_{adj} = \frac{E_{obs} - E_{null}}{E_{max} - E_{null}}$$

where  $E_{obs}$  is the observed BIC/TNW value,  $E_{null}$  is the average BIC/TNW value generated by a Monte Carlo resampling procedure which assumes each individual samples randomly from a shared resource pool (explanation in electronic supplemental materials), and  $E_{max}$  is the maximum possible BIC/TNW value for a given population. The variable  $E_{max}$  can be calculated using the equation 1-1/*P*, where *P* is the number of prey categories (families; see electronic supplemental materials) consumed by a predator population.

inherent temporal and spatial prey variability, we needed to control for these potentially
confounding factors before we could quantify INS. Therefore, we tested each stomach contents
dataset to see if the datasets varied as a function of the variables capture season, capture year,
capture location, size, and sex (Table S1) using principal component analysis and multivariate

153 analysis of variance (MANOVA; Araujo et al. 2007; explanation in electronic supplemental 154 materials). If MANOVAs revealed that any of the independent variables were significant predictors of stomach contents variation in any given dataset, we divided the dataset into smaller 155 subsets to remove the bias (e.g., splitting the dataset into male and female subsets to control for 156 sex differences in diets). We chose the subsets of each dataset with sample sizes of at least 10 157 158 individuals for BIC/TNW analysis using the program IndSpec 1.0 (Bolnick et al. 2002), then converted the BIC/TNW values into  $E_{adj}$  values. We used the program's Monte Carlo procedure 159 to test the null hypothesis that any observed variation in diet was caused by individuals sampling 160 161 randomly from a shared resource pool (Araujo et al. 2007; explanation in electronic 162 supplemental materials). 163

## 164 *Stable isotope collection and analyses*

For the purposes of INS analyses, SIA can be very useful because different tissues within 165 the same consumer can incorporate isotopes from the diet over different time periods (i.e., 166 167 "turnover rates"; Dalerum and Angerbjorn 2005). Thus, multiple tissues collected from one individual can provide insight into the relative stability of dietary patterns – or at least basal 168 169 carbon sources consumed – over multiple timescales. Although identifying specific consumed prey taxa with SIA is difficult for generalist carnivores with broad diets, SIA is still useful 170 because values of  $\delta^{13}$ C are indicative of the origin of a consumer's nutrients (Fry 2006). 171 172 Tissue-specific turnover rates can vary widely between species (Dalerum and Angerbjorn 2005). A diet-switch study of juvenile alligators (Rosenblatt and Heithaus 2013) revealed that the 173 approximate complete turnover time of alligator plasma for  $\delta^{13}$ C (252 days) was roughly half 174 that of red blood cells (RBCs; 566 days). Therefore, if  $\delta^{13}$ C values for an alligator were similar 175

across these tissue types it would suggest that the mixture of carbon pools used across an eight month period prior to sample collection is similar to that used across a 19-month period prior to sample collection. Differing  $\delta^{13}$ C values would indicate shifts in the relative contributions of different carbon pools across these timescales. For our analyses we used stable isotope data from plasma and RBC samples from 214 alligators collected between 2008 and 2012 from six sites (Table 1; see Rosenblatt and Heithaus (2013) for sample collection procedures and the electronic supplemental materials for laboratory analysis procedures).

Before assessing the prevalence of INS in the alligator populations using SIA we needed 183 184 to remove the possibility that our results were affected by variable fractionation factors between the two tissues (Dalerum and Angerbjorn 2005). Therefore, we subtracted experimentally 185 determined fractionation values (isotopic differences between tissues and diet) for each alligator 186 tissue (+0.35‰ for plasma  $\delta^{15}$ N, -0.04‰ for plasma  $\delta^{13}$ C, +0.95‰ for RBC  $\delta^{15}$ N, and +0.03‰ 187 for RBC  $\delta^{13}$ C; Rosenblatt and Heithaus 2013) from the  $\delta$  values for each tissue. Then, 188 MANOVAs were used to determine if the stable isotope datasets were significantly affected by 189 190 the variables capture season, capture location, size, or sex. We only focused on the  $\delta^{13}$ C values 191 of the two tissues as the dependent variables because they contain information about nutrient origins. After controlling for possible confounding variables and the potential effects of 192 fractionation factors, we determined correlation coefficients for each dataset to see if the 193 relationships between the  $\delta^{13}$ C values of plasma and RBCs were linear and positive (i.e., 194 exhibited correlation coefficients close to one). We then employed linear regression to further 195 examine the relationship between the  $\delta^{13}$ C values of plasma and RBCs. If individual alligators 196 exhibited stable foraging patterns over the long time periods represented by the two tissues, we 197 would expect the  $\delta^{13}$ C values of plasma and RBCs to be highly correlated, with linear regression 198

- best-fit lines characterized by high  $R^2$  values and slopes close to one. Conversely, if alligators
- 200 exhibited more variable foraging patterns we would expect best-fit lines characterized by low  $R^2$
- values and slopes farther away from one. All statistical analyses were performed in SPSS 17.0
- 202 (IBM, Armonk, NY, USA) and SigmaPlot 11.0 (Systat, Chicago, IL).

203 **Results** 

#### 204 Stomach contents

The alligators in this study consumed a diverse array of prey. The number of prey 205 families consumed per study ranged from 11 to 38 (mean =  $23.1 \pm 9.2$  SD) and the mean number 206 of individual prey items consumed per alligator per study ranged from 4.5 to 34.8 (mean =  $16.5 \pm$ 207 208 11.3 SD). Prey included crustaceans, mollusks, fishes, amphibians, reptiles, mammals, birds, aquatic and terrestrial insects, and seeds (Table S2). Alligators with empty stomachs made up 209 6.1% (N = 89) of all the individuals (mean per study =  $7.3\% \pm 6.7$  SD), and 27.8% (N = 403) of 210 alligator stomachs contained only 1 prey item or 2 prey items in the same prey category (mean 211 per study =  $26.3\% \pm 14.5$  SD). Therefore, according to our sorting rules, we removed 33.9% of 212 the alligators from further analyses, leaving 960 individuals. 213 The PCA-MANOVA analyses revealed that for most of the datasets (7 out of 12 total) 214 there existed differences in diet separate from INS (Table S1). Capture location was a significant 215 predictor of dietary patterns in 60% (3 of 5) of lake studies and 67% (2 of 3) of estuarine 216 217 habitats. Capture season was a significant predictor of dietary patterns in all three of the studies in which it was included and capture year was a significant predictor in two (67%) of the studies. 218 219 Alligator size was a significant predictor of dietary patterns in 58% of the studies (7 of 12) whereas sex was a significant predictor in two of the nine (22%) studies in which it was 220 included. 221 222 After subdividing each dataset to control for significant predictor variables, the resulting Shark River dataset was removed from further analyses because it was heavily skewed by two 223

individuals that each consumed thousands of anchovies (Engraulidae). The remaining 28

populations and sub-populations only included 23 monophagous individuals between them and

226 produced  $E_{adj}$  specialization values that varied widely, with a minimum of 0.03 and a maximum 227 of 0.74 (mean =  $0.39 \pm 0.19$  SD; Table 2). The Monte Carlo simulations revealed that despite the wide range of observed BIC/TNW values, all but one population exhibited significantly greater 228 229 specialization than predicted by chance (Table 2). The  $E_{adi}$  values were not affected by sample size (linear regression:  $R^2 = 0.007$ , P = 0.7), but the  $E_{adi}$  values were significantly different 230 between habitat types (t-test:  $t_{26} = -2.7$ , P = 0.01), with populations from lakes exhibiting lower 231 232 values (mean =  $0.35 \pm 0.17$  SD) than populations from coastal habitats (mean =  $0.58 \pm 0.14$  SD). Mollusks were one of the most frequently consumed prey groups across the populations 233 and there was a significant negative relationship between the  $E_{adj}$  values of a population and the 234 number of mollusks consumed as a percentage of the population's total diet ( $R^2 = 0.3$ , P = 0.003; 235 Fig. 2). There were also striking differences in mollusk consumption between lake and coastal 236 237 populations: lake populations consumed significantly more mollusks per individual (mean = 9.0 $\pm$  12.0 SD) than coastal populations (mean = 0.2  $\pm$  0.1 SD; Mann-Whitney U test: U = 1, P < 238 (0.001) and significantly more mollusks as a percentage of the population's total diet (mean = 239 240 51.0%  $\pm$  26.0 SD) than coastal populations (mean = 1.5%  $\pm$  2 SD; U = 1, P < 0.001). Lastly, our analyses indicated lake alligator populations used fewer potential prey categories present in their 241 242 habitat (48%  $\pm$  21 SD) when compared to coastal populations (70%  $\pm$  35 SD), though this difference was not statistically significant (Mann-Whitney U test: U = 37, P = 0.2). 243

244

245 Stable isotopes

The fractionation-corrected  $\delta^{13}$ C values from both plasma and RBCs varied in the stable isotope datasets with five of the six populations exhibiting wide  $\delta^{13}$ C ranges (Table 3). We did not use the Lake Woodruff isotopes in our analyses because the  $\delta^{13}$ C ranges for both plasma and RBCs (Table 3) were too small to confidently estimate INS in this population. The large  $\delta^{13}$ C ranges in the other alligator populations are indicative of the large ranges in  $\delta^{13}$ C values of resource pools available to the alligator populations in each habitat (Peterson and Howarth 1987; Gu et al. 1997; Rosenblatt and Heithaus 2011; Adams and Paperno 2012) and that individuals within these populations display considerable variation in the proportions of food they consume from these different resource pools over the timescales of plasma and RBC turnover.

Capture location (freshwater/intermediate/marine habitats) was a significant predictor of 255  $\delta^{13}$ C values in two of the three estuarine studies (Table S1). Also, sex had a significant effect on 256  $\delta^{13}$ C values in the Lake Apopka study so only males were included in analyses (Table S1). In 257 addition, size had a significant effect on  $\delta^{13}$ C values in the Sapelo Island study (Table S1), with 258 the two smaller alligator size classes displaying much lower  $\delta^{13}$ C values than the two larger size 259 classes. For eight out of nine subpopulations from the five isotope datasets plasma and RBC  $\delta^{13}$ C 260 values exhibited tight linear relationships, with correlation coefficient (Pearson's r) values 261 ranging from 0.88 to 0.99 (mean =  $0.96 \pm 0.04$  SD). The Guana River population was the lone 262 outlier with a Pearson's r value of 0.52. The results of linear regression also showed that the  $\delta^{13}$ C 263 values of the two tissues were highly correlated, with  $R^2$  values ranging from 0.77 to 0.99 (mean 264  $= 0.92 \pm 0.07$  SD, all P < 0.001) and with slopes very close to 1, ranging from 0.79 to 1.08 265 (mean =  $0.96 \pm 0.09$  SD; Fig. 3). Indeed, seven of the eight slope values did not differ 266 significantly from a slope of 1 (all P > 0.1). The lone outlier again was the Guana River 267 population, which also showed a positive relationship but a lower  $R^2$  value (0.28) and slope 268 (0.49).269

#### 270 Discussion

Our results show that species thought to be dietary generalists and that exhibit broad 271 population level diets can actually specialize at the individual level. Stomach contents analysis 272 273 revealed that alligator populations exhibit different short-term dietary patterns, ranging from generalized to more specialized populations. We also found the strength of short-term INS was 274 context dependent, with populations from lakes containing many easy-capture prey exhibiting 275 276 lower INS than populations from other habitats containing fewer easy-capture prey. In addition, SIA revealed long-term stability in the use of different carbon pools by individual alligators 277 despite large differences between the  $\delta^{13}$ C values of those carbon pools: eight out of nine 278 populations we studied from both lake and coastal habitats contained individuals with 279 remarkably consistent  $\delta^{13}$ C values over moderate to long time spans, indicating that alligators 280 281 generally maintain the same foraging patterns through time regardless of habitat type. Importantly, INS existed in addition to the dietary variation caused by differences between sexes, 282 life stages, seasons, years, and locations. 283

Consistent with our hypotheses, our findings suggest that one of the main drivers of INS, 284 at least in the short-term, is habitat heterogeneity and prey community composition. Lower  $E_{adi}$ 285 286 values in alligator populations inhabiting lakes could be a consequence of low habitat variation in lakes and their well-defined boundaries. For example, Lake Apopka in central Florida, whose 287 alligator population was sampled several times across our datasets, has been a shallow, 288 289 homogenous, turbid, algal-dominated lake lacking a diverse fishery since 1947 (Bachmann et al. 1999). In contrast, the estuarine habitats sampled across our datasets contain a diversity of 290 aquatic habitats (freshwater, estuarine, marine) and have diffuse boundaries, thus potentially 291 292 allowing for greater movement of alligators and greater ingress of possible prey taxa. In fact,

movement studies indicate that alligators inhabiting lakes move shorter distances and occupy
smaller ranges than alligators in coastal habitats, and lake alligators largely restrict themselves to
littoral areas (Goodwin and Marion 1979; Rosenblatt et al. 2013). Together, these observations
suggest that lake-bound alligators have access to less distinct habitat types and prey groups than
coastal populations. Consistent with this hypothesis, our analyses indicated lake alligator
populations used fewer prey categories present in their habitat when compared to non-lake
populations, though this result was not statistically significant.

Though the specific effects of habitat heterogeneity on INS have not been examined 300 301 extensively, our findings agree with the few studies that have been done. For example, in a study of wolf (*Canis lupus*) feeding patterns in coastal British Columbia researchers found that wolves 302 inhabiting island habitats exhibited relatively more among-individual dietary variation than 303 mainland wolves. This difference was at least partially attributable to the fact that island wolves 304 had access to multiple food webs (terrestrial and marine) embedded in a spatially heterogenous 305 area while mainland wolves only used one food web (terrestrial) in a more spatially homogenous 306 307 area (Darimont et al. 2009). Furthermore, trophic niche width of a population, a factor that can affect INS, also can decrease as habitat heterogeneity decreases (Layman et al. 2007). These 308 309 studies broadly agree with our results, but more research on the effects of habitat heterogeneity on INS across a greater diversity of species and habitats is needed to confirm our findings. 310

Our results further suggest that prey community composition can drive short-term INS because the  $E_{adj}$  values were negatively correlated with the number of mollusks consumed by a population as a percentage of the whole diet. We hypothesize that the presence of abundant easycapture prey like mollusks in certain habitats may allow the majority of alligators in those habitats to take frequent advantage of a resource so easily found and consumed. However, when 316 mollusks are not as abundant alligators must pursue more mobile prev that are harder to capture. 317 These contrasting contexts would make each individual's short-term diet in mollusk-rich habitats (i.e., lakes; Darby et al. 2006) more similar to conspecifics, thereby decreasing INS, but more 318 319 different from conspecifics in habitats containing less abundant mollusks, thereby increasing INS (e.g., Baird et al. 1992; Tinker et al. 2008). In the latter context dietary variation between 320 individuals could be caused by resource patchiness rather than behavioral specialization, but 321 available evidence suggests that alligators do not move randomly through their environment 322 because individuals visit the same foraging areas over multiple years (Rosenblatt and Heithaus 323 324 2011; Rosenblatt et al. 2013). Although individuals may return to similar locations for periods of time, their foraging trips take them to areas with prey supported by different carbon sources (e.g. 325 marine versus freshwater production). Thus, INS in habitats dominated by more mobile prey 326 327 could be caused by fine-scale associations between individual alligators and specific areas or by individuals learning to eat different prey. 328

The effects of prey capture difficulty on INS have not been extensively studied, but our 329 330 results agree with those from killer whales which show that as prey capture difficulty increases INS increases as well (Baird et al. 1992; Ford et al. 1998). However, a more recent study on 331 332 much smaller predators (wasps; Crabronidae and Sphecidae) showed the opposite effect of prey capture difficulty on INS (Polidori et al. 2013). These incongruent results may be the result of 333 different habitat types (aquatic vs. terrestrial), differences in prey modes of movement (flying vs. 334 335 swimming), or differences in scale. Regardless of the cause of the differences it is clear that the effects of prey capture difficulty can affect INS and more research is needed to identify the 336 mechanisms involved. 337

Stomach contents analysis has specific drawbacks pertaining to INS. For example, SCA 338 can be biased by variable digestive rates of different prey items. Prey with indigestible parts 339 (e.g., arthropods with chitinous exoskeletons, turtles, mollusks) may become over-represented in 340 the diet because they remain in the stomach longer than soft-bodied vertebrate prey (Garnett 341 342 1985; Janes and Gutzke 2002; Nifong et al. 2012). This factor can cause some prey to appear 343 more frequently and in higher numbers in stomachs, which may bias dietary analyses. However, most of the prey groups consumed by alligators in our synthesis have at least one indigestible 344 part (e.g., mammal hair, bird feathers, crustacean exoskeletons, snail shells) which means 345 346 variable digestive rates likely did not have a large effect on our results, with the exception of a possible underestimation of amphibians and fishes (Table S2). 347

There are three additional caveats associated with our stomach contents analyses. First, 348 we are confident that alligators exhibit non-random foraging behaviors, thereby violating one of 349 the assumptions of the Monte Carlo null hypothesis testing procedure. This means that the 350 statistical significance of our INS results is not conservative and should be viewed with caution. 351 352 Second, intraspecific competition is known to affect INS (Estes et al. 2003; Tinker et al. 2008) 353 and to account for this factor we would need to include alligator density and prey density in our 354 analyses. Unfortunately, both of these forms of data were not available for any of the locations we included in our analyses, so there is a possibility that our INS results were also affected by 355 differences in intraspecific competition between habitat types. Third, we grouped alligator prey 356 357 by family because we assumed that alligators do not discriminate between prey on the species or even genus level. If alligators do actually show preferences on the species or genus level, then 358 359 our INS estimates would be conservative and real alligator INS values may be higher, i.e. 360 alligator populations may be more specialized. In contrast, if the opposite were true and

alligators do not discriminate between prey even on the family level, then alligator populationsmay actually be less specialized.

SIA revealed that alligators generally exhibited highly consistent use of different carbon 363 pools across longer time frames. The consistent  $\delta^{13}$ C values we observed across tissues could 364 have been caused by specialization on specific prey but could also be attributed to consistent use 365 of specific habitats and movement patterns. We hypothesize that the latter explanation is more 366 likely, i.e. that alligators specialize in certain behavioral patterns, like specific foraging and 367 movement tactics. This possibility was partially accounted for in our study by separating 368 369 analyses for individuals captured in different habitats, but because of their high mobility 370 alligators captured in the same habitat still could move across the landscape in different ways. Our hypothesis is supported by a previous study of alligator movement patterns in the Shark 371 River where alligators exhibited temporally stable variation in movement patterns (Rosenblatt 372 and Heithaus 2011; Rosenblatt et al. 2013): half of the alligators regularly commuted between 373 freshwater/estuarine and marine habitats and the stable isotope values of their skin indicated that 374 375 they were consistently feeding in two different food webs, whereas other alligators limited their movements and feeding strictly to freshwater/estuarine habitats. Similar patterns appear to occur 376 377 in sea turtles, although on a different time scale, in that individual turtles will use consistent "corridors" to move between nesting and foraging areas and will return to the same foraging 378 areas repeatedly, or individuals from the same nesting populations may be either coastal or 379 380 pelagic in their foraging habitats (see Heithaus 2013 for a review). Our study and others point to the difficulties in using SIA to infer INS behaviors as a standalone data source when turnover 381 rates of even "fast" tissues are long, as in most reptiles (Rosenblatt and Heithaus 2013), because 382 383 in such situations SIA fails to capture short-term dietary choices. Therefore, we echo previous

studies in suggesting that INS studies should combine isotope data with other behavioral data where possible. We should also note that the turnover rate of alligator plasma is almost 50% of the turnover rate for alligator RBCs, thus inherently biasing our analyses towards strong correlations between the  $\delta^{13}$ C values of the two tissues. However, despite this methodological artifact we are confident in our results because of the remarkable strength of the correlations across almost every alligator population we sampled.

Many current ecosystem management and conservation strategies assume that all apex 390 predators in a population will have similar effects on prey populations. However, if behaviors 391 392 vary consistently across individuals, as our study and those of other taxa (Bolnick et al. 2003) suggest, then one-size-fits-all conservation and management strategies may have unintended 393 consequences in many systems. For example, the Shark River is part of the Everglades which is 394 an ecosystem currently undergoing large-scale restoration (Doren et al. 2009). Restoration 395 activities are expected to bring more freshwater to the area, thereby decreasing salinity and 396 potentially increasing habitat quality for alligators that are dependent on freshwater (Mazzotti 397 398 and Brandt 1994). Since many of the alligators in this system appear to specialize in exploiting 399 the marine food web seasonally (Rosenblatt and Heithaus 2011), increased freshwater flow may 400 positively impact this group by allowing them to access marine resources for a larger portion of each year (though the distribution of marine prey may change as well). However, individuals that 401 do not currently take advantage of marine resources may not experience the same future benefits. 402 403 In addition, the spatial scale of alligator movements, and therefore potential coupling of food webs and nutrient transport (Rosenblatt and Heithaus 2011), likely will change. As freshwater 404 405 moves seaward, any individuals exploiting marine food webs will not need to move as far upstream for osmoregulation. Similarly, in northern Kenya, African wild dog (Lycaon pictus) 406

populations outside of protected areas exhibit specialized feeding on smaller prey than
populations inside protected areas (Woodroffe et al. 2007). Thus, if wild dog conservation efforts
were solely aimed at bolstering large prey populations some wild dogs may see no benefit unless
their feeding patterns are very plastic. Different restoration and conservation scenarios, therefore,
may differentially benefit subsets of apex predator populations. As a result, restoration and
conservation actions may not achieve their intended goals if individual behavioral specialization
is not explicitly considered during the planning process.

Our results also have important implications for food web theory and ecosystem models. 414 415 Theoretical analyses have shown that top predators that couple different food webs through their 416 foraging and movement behaviors contribute both local and non-local stability to the food webs (Rooney et al. 2006), but such models treat individuals in predator populations as behaviorally 417 homogenous. By incorporating behaviorally heterogenous predator populations into food web 418 and ecosystem models, realism of theoretical analyses and scenarios will increase (Quevedo et 419 al. 2009). These next-generation models may reveal insights into the potentially nuanced roles of 420 421 top predators in ecosystems and allow for the development of additional testable hypotheses.

Ultimately, our research shows that INS in large apex predators can vary substantially 422 423 among populations and the degree of inter-individual variation is likely affected by habitat type and prey community composition. Future research could make valuable contributions to our 424 understanding of INS by investigating the relative effects of prey community composition, niche 425 426 width, and inter- and intraspecific competition across a wider range of taxonomic groups, body sizes, and trophic levels. In the context of INS in large apex predators, it remains to be seen how 427 428 subpopulations exhibiting different behaviors will respond to specific ecosystem conservation 429 and management scenarios, but it is clear that INS needs to be explicitly considered in such plans 430 as well as in food web models.

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Data source	Locations	Habitat type	Duration	Collection method	Ν
	Sto	omach contents			
Delany and Abercrombie 1986	andOrange, Lochloosa, & Newnansombie 1986Lakes, FL		1981-1983	Hunter harvested alligators	349
Delany et al. 1988	Duval, St. Johns, Alachua, Marion, Citrus, & Lake counties, FL	Lake	1977	1977 Sacrificed nuisance alligators	
Delany 1990	Orange Lake, FL	Lake	1986	Sacrificed alligators	77
Elsey et al. 1992	Marsh Island, LA	Island	1991	Hunter harvested alligators	101
Delany et al. 1999	al. 1999 Rodman, George, Hancock, & Lake 1985 Hunter harvested alligators		231		
Rice et al. 2007	Apopka, Griffin, & Woodruff Lakes, FL	Lake	2001-2003	Hose-Heimlich stomach flushing	172
Gabrey 2010	Lafourche, Terrebonne, Cameron, Vermilion, & St. Charles parishes, LA	Marsh	2002-2004	Hunter harvested alligators	212
	Apopka & Woodruff Lakes, FL	Lake	2010	Hose-Heimlich stomach flushing and necropsies	29
This should	Merritt Island, FL	Island	2010	Hose-Heimlich stomach flushing and necropsies	10
This study	Shark River, FL	Estuary	2009-2011	Hose-Heimlich stomach flushing	54
	Guana River, FL	Estuary	2011-2012	Hose-Heimlich stomach flushing	40
J. Nifong (unpub. data)	Sapelo Island, GA	Island	2007-2010	Hose-Heimlich stomach flushing	99
	S	table isotopes			
	Apopka & Woodruff Lakes, FL	Lake	2010	NA	29
This study	Merritt Island, FL	Island	2010	NA	10
rnis study	Shark River, FL	Estuary	2008-2011	NA	79
	Guana River, FL	Estuary	2011-2012	NA	40
J. Nifong (unpub. data)	Sapelo Island, GA	Island	2009-2010	NA	56

Table 1 Summary information for the American alligator stomach contents and stable isotope studies used in analyses.

Table 2 Results of American alligator stomach contents specialization analyses. BIC/TNW = Between Individual Component/Total Niche Width
 (observed), BIC/TNW MC = mean value generated by Monte Carlo simulations, TL = Total Length of alligators, SVL = Snout-Vent Length of
 alligators.

Data source	Location	Sample date	Sex	Size (TL; cm)	N	BIC/TNW	BIC/TNW MC	E <sub>max</sub> (prey	Eadjusted
	Location				IN	(E <sub>obs</sub> )	(E <sub>null</sub> )	categories)	
	Orange Lake, FL	Fall 1983	M <i>,</i> F	200-250	22	0.71**	0.20	0.94 (17)	0.69
	Lochloosa Lake, FL	Fall 1981	M, F	150-200	10	0.68**	0.20	0.9 (10)	0.69
	Orange Lake, FL	Fall 1982	M, F	150-200	22	0.58**	0.38	0.93 (14)	0.36
	Orange Lake, FL	Fall 1981	M, F	250-300	10	0.55**	0.40	0.94 (16)	0.28
Dolony and	Orange Lake, FL	Fall 1983	M, F	100-150	12	0.52**	0.28	0.93 (14)	0.37
Aborcrombio 1086	Orange Lake, FL	Fall 1982	M, F	200-250	19	0.52**	0.42	0.93 (15)	0.19
ADELCIONDIE 1980	Orange Lake, FL	Fall 1982	M, F	100-150	13	0.52**	0.38	0.92 (13)	0.26
	Orange Lake, FL	Fall 1983	M, F	250-300	13	0.51**	0.32	0.92 (13)	0.32
	Orange Lake, FL	Fall 1982	M, F	250-300	10	0.48**	0.35	0.88 (8)	0.25
	Newnan's Lake, FL	Fall 1983	М	300-350	15	0.48**	0.32	0.91 (11)	0.27
	Newnan's Lake, FL	Fall 1983	M, F	200-250	11	0.43**	0.22	0.9 (10)	0.31
Delany et al. 1988	Griffin & Tsala Apopka Lakes, FL	Summer 1977	M, F	100-350	27	0.58	0.57	0.93 (14)	0.03
	Orange Lake, FL	Fall 1986	NA	50-100	27	0.56**	0.25	0.91 (11)	0.47
Delany 1990	Orange Lake, FL	Fall 1986	NA	0-50	27	0.54**	0.24	0.9 (10)	0.45
	Orange Lake, FL	Fall 1986	NA	100-150	13	0.48**	0.18	0.92 (13)	0.4
Elsey et al. 1992	Marsh Island, LA	Summer 1991	M, F	100-200	81	0.57**	0.22	0.95 (22)	0.48
	Rodman Lake, FL	Summer 1985	М	250-300	15	0.45**	0.11	0.94 (16)	0.41
	George Lake, FL	Summer 1985	M, F	150-200	18	0.46**	0.18	0.92 (12)	0.38
Delany and Abercrombie 1986 Delany and Abercrombie 1986 Delany et al. 1988 Delany et al. 1999 Delany et al. 1990 Delany e	Summer 1985	M, F	100-150	12	0.44**	0.16	0.89 (9)	0.38	
Delatty et al. 1999	Rodman Lake, FL	Summer 1985	M, F	200-250	11	0.33**	0.08	0.89 (9)	0.31
	George Lake, FL	Summer 1985	M, F	200-250	10	0.55**	0.44	0.89 (9)	0.25
	Rodman Lake, FL	Summer 1985	М	300-350	10	0.24**	0.14	0.91 (11)	0.13
Rice et al. 2007	Woodruff Lake, FL	Fall 2002	M, F	200-250	12	0.70**	0.10	0.92 (12)	0.73
Gabrey 2010	Freshwater marsh, LA	Fall 2004	М	75-100 (SVL)	15	0.79**	0.57	0.9 (10)	0.67
	Merritt Island, FL	Spring 2010	М	250-350	7	0.69**	0.07	0.91 (11)	0.74
This study	Guana River, FL	Summer 2011-2012	M, F	50-250	30	0.64**	0.11	0.95 (21)	0.63
	Apopka & Woodruff Lakes, FL	Spring 2010	M, F	200-350	20	0.47**	0.35	0.92 (13)	0.21
J. Nifong (unpub. data)	Sapelo Island, GA	Spring 2008	М	50-100	10	0.50**	0.29	0.83 (6)	0.39

572 \*\**P* < 0.001 (Monte Carlo bootstraps, 500 simulations)

573 All others, *P* = 0.3

571

574 Table 3 Summary of fractionation-corrected  $\delta^{13}$ C ranges for plasma and red blood cells (RBC) for each American alligator population used in the 575 study.

Location	Min plasma δ <sup>13</sup> C (‰)	Max plasma δ <sup>13</sup> C (‰)	Plasma δ <sup>13</sup> C mean ± SD (‰)	Min RBC δ <sup>13</sup> C (‰)	Max RBC δ <sup>13</sup> C (‰)	RBC $\delta^{13}$ C mean ± SD (‰)
Sapelo Island, GA	-29.4	-15.3	-21.5 ± 4.5	-28.0	-14.0	-21.0 ± 4.5
Lake Apopka, FL	-25.9	-11.9	-16.5 ± 4.9	-25.2	-12.1	-16.3 ± 4.8
Shark River, FL	-30.3	-19.2	-26.2 ± 2.1	-28.7	-21.1	-26.0 ± 1.7
Guana River, FL	-24.6	-18.0	-20.8 ± 1.3	-25.0	-18.0	-21.1 ± 1.4
Merritt Island, FL	-22.8	-16.8	-19.0 ± 2.0	-23.2	-15.2	-18.0 ± 2.3
Lake Woodruff, FL	-26.4	-24.4	$-25.4 \pm 0.6$	-26.0	-23.3	-24.8 ± 0.7

576

577

# 578 Figure captions

579

Fig. 1 Map of alligator sampling locations used for diet specialization analyses. Circles = lakes,
 squares = islands, triangles = marshes, and diamonds = rivers/estuaries. Black shapes represent
 locations where only stomach contents were collected and gray shapes represent sites where both
 stomach contents and tissues for stable isotope analysis were collected

- 584
- **Fig. 2** Specialization values  $(E_{adj})$  of different American alligator populations as predicted by
- mollusk consumption (total number of mollusks consumed by population divided by total
- number of all prey items consumed by population). Black line is linear regression trend line
- 588
- **Fig. 3** Linear regression plots depicting relationships between alligator plasma  $\delta^{13}$ C values and
- red blood cell  $\delta^{13}$ C values. **a** Lake Apopka, FL (males, lake habitat, 250-400 cm total length
- (TL)), **b** Sapelo Island, GA (males and females, marine habitat, 100-150 cm TL), **c** Sapelo
- Island, GA (males, marine habitat, 50-100 cm TL), **d** Shark River, FL (males and females,
- estuarine habitat, 150-300 cm TL). All P < 0.001. Note different scales on axes