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# Multi-Tissue Stable Isotope Analysis and Acoustic Telemetry Reveal Seasonal Variability in the Trophic Interactions of Juvenile Bull Sharks in a Coastal Estuary

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1 **Multi-tissue stable isotope analysis and acoustic telemetry reveal seasonal variability in the**  
2 **trophic interactions of juvenile bull sharks in a coastal estuary**

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8 Running head: seasonal changes in bull shark diets

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11

12 **Summary**

- 13 1. Understanding how natural and anthropogenic drivers affect extant food webs is critical  
14 to predicting the impacts of climate change and habitat alterations on ecosystem  
15 dynamics.
- 16 2. In the Florida Everglades, seasonal reductions in freshwater flow and precipitation lead to  
17 annual migrations of aquatic taxa from marsh habitats to deep-water refugia in estuaries.  
18 The timing and intensity of freshwater reductions, however, will be modified by ongoing  
19 ecosystem restoration and predicted climate change.
- 20 3. Understanding the importance of seasonally pulsed resources to predators is critical to  
21 predicting the impacts of management and climate change on their populations. As with  
22 many large predators, however, it is difficult to determine to what extent predators like  
23 bull sharks (*Carcharhinus leucas*) in the coastal Everglades make use of prey pulses  
24 currently.
- 25 4. We used passive acoustic telemetry to determine whether shark movements responded to  
26 the pulse of marsh prey. To investigate the possibility that sharks fed on marsh prey, we  
27 modeled the predicted dynamics of stable isotope values in bull shark blood and plasma  
28 under different assumptions of temporal variability in shark diets and physiological  
29 dynamics of tissue turnover and isotopic discrimination.
- 30 5. Bull sharks increased their use of upstream channels during the late dry season, and  
31 although our previous work shows long-term specialization in the diets of sharks, stable  
32 isotope values suggested that some individuals adjusted their diets to take advantage of  
33 prey entering the system from the marsh, and as such this may be an important resource  
34 for the nursery.

- 35 6. Restoration efforts are predicted to increase hydroperiods and marsh water levels, likely  
36 shifting the timing, duration, and intensity of prey pulses, which could have negative  
37 consequences for the bull shark population and/or induce shifts in behavior.
- 38 7. Understanding the factors influencing the propensity to specialize or adopt more flexible  
39 trophic interactions will be an important step in fully understanding the ecological role of  
40 predators and how ecological roles may vary with environmental and anthropogenic  
41 changes.

42

43 **Key words:**  $\delta^{13}\text{C}$ , food webs, habitat use, movements, shark nursery, trophic ecology

44

45 **Introduction**

46 Coastal ecosystems experience considerable daily and seasonal variation in  
47 environmental conditions (Lewis 2001; Kennish 2002). Also, they have been, and continue to  
48 be, heavily influenced by human activities that have contributed to shifts in community  
49 composition and have potentially altered the ecological roles of species (e.g. Cloern 2001;  
50 Jackson et al. 2001; Parmesan & Yohe 2003). Within coastal ecosystems, predators serve  
51 important roles in controlling prey populations, linking disparate food webs, and transporting  
52 biomass and nutrients across habitat boundaries (e.g. Bowen 1997; Darimont, Paquet &  
53 Reimchen 2009). Thus, understanding how predators are affected by temporally variable and  
54 ephemeral food sources is important for understanding the trophic dynamics of a system.  
55 However, this can be challenging, because predators are often highly mobile with relatively large  
56 home ranges. Consequently, manipulative studies can be difficult to execute and/or lead to  
57 biased results. Data quantifying behavioral variability in response to natural variation in food  
58 sources can provide valuable insight in the roles predators play. In addition, there is a growing  
59 need to understand how both natural and anthropogenic factors influence variability in trophic  
60 interactions to predict how they may affect the ecological roles of species and ultimately  
61 ecosystem dynamics.

62 Seasonal changes in temperature, precipitation, and freshwater flow lead to noticeable  
63 variation in the distribution, abundance, and behavior of many resident and migratory species in  
64 the Florida Coastal Everglades (Chick, Ruetz & Trexler 2004; Ruetz, Trexler & Jordan 2005;  
65 Rehage & Trexler 2006). Therefore, trophic interactions are likely to vary in space and time as  
66 predators and prey move to stay within suitable environmental conditions or to take advantage of  
67 seasonal pulses of prey. These seasonal pulses of prey occur in the coastal Everglades when

68 water levels in freshwater marshes drop and numerous aquatic taxa are forced into deep-water  
69 channels (Rehage & Trexler 2006; Rehage & Loftus 2007). The magnitude and timing of these  
70 pulses are likely to be affected by ecosystem restoration. Freshwater flow is predicted to  
71 increase through freshwater marshes, likely reducing the duration and intensity of marsh dry-  
72 down (Sklar et al. 2001; Perry 2004; CERP 2006), and therefore the magnitude and timing of  
73 resource pulses into creeks. Thus, understanding the value of this resource pulse in the trophic  
74 ecology of estuarine predators will be important for predicting the consequences of restoration  
75 efforts within the ecosystem.

76 Stable isotope analysis has become a popular method used in ecological studies of food  
77 webs to investigate trophic interactions (reviewed by Layman et al. 2012). Because the materials  
78 eaten by an animal are not immediately incorporated into its tissues, stable isotope values  
79 provide dietary data over a previous timeframe based on the isotopic turnover rate of the sampled  
80 tissue(s) (Gannes, O'Brien & Martinez del Rio 1997; Post 2002; Martinez del Rio et al. 2009).  
81 This lag time can provide a means to investigate the temporal variability in the diet of an  
82 organism by serially sampling parts of metabolically inert tissues or comparing the isotopic  
83 values of multiple tissues with different turnover rates (Bearhop et al. 2004).

84 Metabolically inert tissues, like vibrissae in California sea otters (*Enhydra lutris nereis*),  
85 provide dietary information about a particular time period or event in the life of an organism, and  
86 sequentially sampling inert tissues can provide a dietary record for an organism over its lifetime  
87 (Newsome et al. 2009). Unfortunately, many animals do not have easily accessible tissues that  
88 can be used for serial sampling. One alternative to sequentially sampling metabolically inert  
89 tissues is to sample metabolically active tissue(s) from animals over multiple time periods and  
90 quantify the variability in isotope values over time (Bearhop et al. 2004). This can be an

91 effective method when individuals are easily recaptured (e.g. Drago et al. 2010). But many  
92 ecosystems are open and animals can move across habitat boundaries, making it difficult to  
93 regularly sample the same individuals over time, and limiting the use of this approach.

94 A third strategy to investigate temporal change in diets is to compare the isotopic values  
95 of metabolically active tissues with significantly different turnover rates collected during one  
96 sampling event. A tissue with a fast isotopic turnover rate like blood in Japanese quail (*Coturnix*  
97 *japonica*,  $\delta^{13}\text{C}$  half-life = 11.4 days), provides information on recent trophic interactions, while a  
98 tissue with a slower turnover rate like bone collagen ( $\delta^{13}\text{C}$  half-life = 173.3 days in *C. japonica*),  
99 provides a view of the average trophic interactions over an extended time period (Hobson &  
100 Clark 1992). If the isotope values of a fast turnover tissue are compared to the isotope values of  
101 a slow turnover tissue, the difference in isotope values can provide insight into the temporal  
102 variability of an organism's trophic interactions. Comparing the isotope values of multiple  
103 metabolically active tissues, however, must be conducted carefully because the values of stable  
104 isotopes in the tissues of a consumer are not identical to that of their food due to isotopic  
105 discrimination (Gannes, O'Brien & Martinez del Rio 1997), and different tissues from the same  
106 organism can have different discrimination factors (e.g. Vander Zanden & Rasmussen 2001;  
107 Sweeting et al. 2007; Buchheister & Latour 2010). Thus, understanding the isotopic  
108 discrimination values of the tissues being used is important when investigating temporal  
109 variability in trophic interactions.

110 While stable isotope analysis provides useful information on the trophic interactions of  
111 animals, data from complimentary approaches strengthen inferences about the trophic ecology of  
112 individuals and populations. Acoustic telemetry, for example, provides data on the movements  
113 of tagged animals, and when paired with stable isotope analysis, can provide a powerful tool for

114 elucidating individual- and population-level patterns linking habitat use and diet (e.g.  
115 Papastamatiou et al. 2010; Rosenblatt & Heithaus 2011; Speed et al. 2012). Here, we used a  
116 combination of long-term, passive acoustic tracking, and stable isotopic analysis and modeling  
117 using blood plasma (faster turnover tissue) and whole blood (slower turnover tissue)  $\delta^{13}\text{C}$  values  
118 to investigate whether juvenile bull sharks make use of seasonal prey pulses in the coastal  
119 Everglades.

120

## 121 **Methods**

### 122 *Study species and system*

123 Bull sharks (*Carcharhinus leucas*; Müller & Henle 1839) inhabit coastal and estuarine  
124 waters of the tropics and subtropics around the world, and use coastal estuaries as nurseries  
125 during early years before moving into coastal ocean habitats (Wiley & Simpfendorfer 2007,  
126 Grubbs 2010). Bull sharks can travel between fresh and marine waters with minimal metabolic  
127 costs, and young individuals can be found in salinities ranging from 0.2-41.7 parts per thousand  
128 (Anderson et al. 2006; Steiner, Michel & O'Donnell 2007; Heupel & Simpfendorfer 2008). As a  
129 result, bull sharks can take advantage of a variety of prey types, including teleosts, crustaceans,  
130 cephalopods, and other elasmobranchs in marine, brackish, and freshwater habitats (Snelson &  
131 Williams 1981; Snelson, Mulligan & Williams 1984; O'Connell et al. 2007).

132 The Shark River Estuary of Everglades National Park, Florida, USA (Fig. 1) is primarily  
133 a braided stream system lined by mangroves that extends more than 30 km upstream from the  
134 Gulf of Mexico. The estuary serves as a nursery for juvenile bull sharks year-round, which are  
135 found throughout the entire system (Wiley & Simpfendorfer 2007; Heithaus et al. 2009; Matich  
136 & Heithaus 2012). Seasonal changes in precipitation and freshwater flow lead to noticeable



137 variation in the distribution, abundance, and behavior of many resident and migratory species in  
138 the Florida Coastal Everglades, including the Shark River Estuary (Chick, Ruetz & Trexler 2004;  
139 Ruetz, Trexler & Jordan 2005; Rehage & Trexler 2006). Therefore, trophic interactions vary in  
140 space and time annually as predators and prey move to stay within acceptable environmental  
141 conditions and/or to take advantage of seasonal pulses of prey. Seasonal pulses of freshwater  
142 prey into mangrove-lined creeks in the upstream region of the Shark River Estuary occur when  
143 water levels in freshwater marshes drop during the dry season (Rehage & Trexler 2006; Rehage  
144 & Loftus 2007; Fig. 1), and teleost predators rely on this prey pulse as an important seasonal  
145 component of their diets (Boucek & Rehage *in press*). Our previous work revealed that juvenile  
146 bull sharks have relatively high levels of individual dietary specialization in the Shark River  
147 Estuary (Matich, Heithaus & Layman 2011). Yet, stable isotope analysis revealed that some  
148 individuals (*ca.* 13%) exhibit temporal variability in their trophic interactions, possibly driven by  
149 use of this seasonal pulse of marsh prey.

150 Marsh water levels serve as a seasonal indicator for when taxa migrate from the marsh  
151 into deep-water refuges. For the purposes of our analyses, we used water level data from United  
152 States Geological Survey water station 252820080505400 Everglades National Park  
153 (N25°28'20", W80°50'54"; Fig. 1) adjacent to our study system. When marsh water elevation  
154 drops below 10 cm in depth in reference to elevation, the marsh becomes unsuitable for large  
155 aquatic taxa (> 8 cm), which are forced to seek out deep-water habitat. As such, the dry season,  
156 in reference to water levels, occurs when marsh water elevations are less than 10 cm and the wet  
157 season occurs when water levels are greater than 10 cm. These thresholds have been used in  
158 studies of movements of Everglades marsh taxa into estuarine creeks (e.g. Chick, Ruetz &  
159 Trexler 2004; Rehage & Loftus 2007; Parkos, Ruetz & Trexler 2011). Because the abundance of

160 marsh prey within mangrove-lined creeks changes considerably within seasons (Rehage &  
161 Loftus 2007), we further divided each season into sub-seasons (i.e. early and late dry seasons,  
162 and early and late wet seasons). During our study, the late wet season of 2008/2009 ended on 29  
163 Feb 2009, and the early dry season was from 1 Mar to 13 Apr 2009. The late dry season was  
164 from 14 Apr to 28 May 2009, the early wet season was from 29 May to 16 Oct 2009, and the late  
165 wet season began 17 Oct 2009.

166 Within the confines of the Shark River Estuary, there are two isotopically distinct food  
167 webs - freshwater/estuarine ( $\delta^{13}\text{C} < -25\text{‰}$ ) and marine ( $\delta^{13}\text{C} > -19\text{‰}$ ; Fry & Smith 2002; Chasar  
168 et al. 2005; Williams & Trexler 2006; Fig. 2). Marsh taxa that enter the estuary during the dry  
169 season prey pulse have more depleted  $\delta^{13}\text{C}$  values (mean  $\pm$  SE =  $-30.5 \pm 0.5\text{‰}$ ) than resident  
170 freshwater/estuarine taxa (mean  $\pm$  SE =  $-28.0 \pm 0.5\text{‰}$ ; Matich & Boucek unpublished data).  
171 These differences in the  $\delta^{13}\text{C}$  values of potential prey species enabled us to investigate seasonal  
172 shifts in bull shark diets between prey with different basal carbon sources, and the potential use  
173 of the freshwater prey pulse, by quantifying temporal variability in the  $\delta^{13}\text{C}$  values of bull shark  
174 tissues.

175

### 176 *Field Sampling*

177 Bull sharks were captured from Oct 2008 to Mar 2012 on ~500 m longlines, fitted with  
178 40-55 14/0 or 15/0 Mustad tuna circle hooks baited with mullet (*Mugil* sp.) and attached to ~2 m  
179 of 400 kg monofilament line (see Heithaus et al. 2009 for further details of sampling equipment).  
180 Captured sharks were processed alongside the sampling vessel, or within a water-filled, aerated  
181 cooler on board. Shark total length was measured to the nearest centimeter. An 18 gauge needle  
182 was used to collect 4 mL of blood from the caudal vein. During collection, 3 mL of blood was

183 placed into BD Vacutainer blood collection vials with neither additives nor interior coating, and  
184 then immediately separated into its components, including plasma, using a centrifuge spun for  
185 one minute at 3000 rpm. The remaining 1 mL of blood was retained in its original composition  
186 (whole blood, “blood” hereafter). Based on several lab studies, plasma has an isotopic half-life  
187 of ~32 days in elasmobranchs (Kim et al. 2012), and likely serves as a short-term diet indicator  
188 for juvenile bull sharks, while blood has an isotopic half-life of ~61 days (MacNeil, Drouillard &  
189 Fisk 2006), and likely serves as a longer-term diet indicator for juvenile bull sharks. As such,  
190 when more dynamic plasma  $\delta^{13}\text{C}$  values are compared to more stable blood  $\delta^{13}\text{C}$  values they can  
191 be used to study short-term changes in the diets of sharks, and provide diet information over the  
192 time-frame juvenile bull sharks may respond to the pulse of marsh prey into the Shark River  
193 Estuary. Importantly, such inter-tissue comparisons are useful even if tissues do not reach full  
194 dietary equilibrium (i.e. four half lives), because they can provide data on the direction (i.e. an  
195 increase of decrease in  $\delta^{13}\text{C}$  depleted prey) and magnitude of dietary change. Tissue samples  
196 were put on ice and frozen before laboratory preparations. All samples were dried and  
197 homogenized prior to stable isotopic analysis.

198       Muscle tissue was collected from known estuarine (*Lutjanus griseus*, *Mugil cephalus*)  
199 and marsh teleosts (*Lepomis marginatus*, *L. microlophus*, *L. punctatus*) that may serve as prey  
200 for juvenile bull sharks (based on gape size of sharks, size of teleosts, and stomach contents  
201 analysis of juvenile bull sharks in other systems: Snelson & Williams 1981; Snelson, Mulligan &  
202 Williams 1984; O’Connell et al. 2007). Samples were collected during ongoing community  
203 level surveys using electrofishing, which occurred during the bull shark study (see Rehage &  
204 Loftus 2007 for description of sampling method). Samples were frozen until being dried and  
205 homogenized in the lab. Stable isotope data from teleosts provided the framework for the

206 difference in  $\delta^{13}\text{C}$  values of estuarine and freshwater prey for stable isotope diet change models  
207 (see below).

208         Passive acoustic tracking was used to quantify the movement patterns of individual bull  
209 sharks to assess their use of upstream areas of the estuary in response to the marsh prey pulse.  
210 From Oct 2008 to May 2009, sharks caught in excellent condition (swimming strongly upon  
211 capture;  $n = 23$ ) were surgically fitted with a Vemco V16-4H transmitter (Vemco, Halifax, NS).  
212 Transmitters were set to emit a unique series of pulses for each shark at a random interval  
213 between 30-90 sec (mean emission interval = 60 sec). Movements of acoustically tagged sharks  
214 were tracked within an array of 43 Vemco VR2 and VR2W acoustic receivers (Fig. 1) from Oct  
215 2008 to Nov 2009. In situ measurements revealed mean detection ranges of receivers were ~500  
216 m (see Rosenblatt & Heithaus 2011 for detection ranges of individual receivers). Each receiver  
217 was attached to a PVC pipe set in a 10 kg cement anchor. Data from receivers were downloaded  
218 every 3-4 months for the duration of the study, and batteries were replaced as needed.

219

#### 220 *Stable isotope analysis*

221         All shark samples were analyzed at the Florida International University Stable Isotope  
222 Facility (29 blood samples and 30 plasma samples) or the Yale Earth System Center for Stable  
223 Isotopic Studies (61 blood samples and 60 plasma samples). Lipid extraction was not performed  
224 because C:N ratios (mean blood =  $2.63 \pm 0.25$  SD; mean plasma =  $2.03 \pm 0.26$  SD) were below  
225 those suggested for extraction or mathematical correction (3.5; Post et al. 2007). To verify  
226 analytical consistency, we randomly selected samples to be analyzed at both Florida  
227 International University and Yale University, for which the variation between resulting  $\delta^{13}\text{C}$   
228 values and  $\delta^{15}\text{N}$  values were  $0.13\text{‰} \pm 0.20$  SE. The standard deviations of standards run for

229 Yale were 0.14‰ for  $\delta^{13}\text{C}$  and 0.22‰ for  $\delta^{15}\text{N}$ , and 0.29‰ for  $\delta^{13}\text{C}$  and 0.24‰ for  $\delta^{15}\text{N}$  for  
230 Florida International. All teleost samples were analyzed at the Florida International University  
231 Stable Isotope Facility.

232

### 233 *Quantitative Analysis*

#### 234 Acoustic tracking

235 We quantified the proportion of days each tagged shark was detected in the upstream  
236 region based on receiver detections of tagged sharks (Fig. 1). We predicted that if sharks fed  
237 from the prey pulse, they would have increased their use of the upstream region where  
238 freshwater taxa enter the system early in the dry season, and decreased their use of the upstream  
239 region later in the dry season when the abundance of marsh prey decreased (Rehage & Loftus  
240 2007). Sharks were only used for analysis if they were present within the array for > 3 months,  
241 and were within the array during the 2009 dry season when the marsh prey pulse was expected to  
242 enter the estuary (1 Mar to 28 May). We used a random effects GLMM to test the effect of  
243 month on the average proportion of days individual sharks were detected by upstream receivers,  
244 with individual as a random effect, and used a Post hoc Tukey's test to test for significant  
245 differences across months. We used linear regression to examine the relationship between marsh  
246 water level and the proportion of sharks detected per day by upstream receivers during the dry  
247 season. Finally, based on the movements of marsh taxa during the dry season, we used t-tests to  
248 investigate if there was a significant change in the use of the upstream region by sharks when  
249 water elevations were  $\leq 0$  cm, between 0-5 cm, and between 0-10 cm to develop predictions for  
250 our diet change models (see below). Investigating shark habitat use in relation to these water  
251 depths allowed us to examine if sharks changed their movement behavior in response to the entry

252 of larger marsh taxa (> 8 cm; enter the estuary from the marsh when water levels drop below 10  
253 cm) or smaller marsh taxa (< 8 cm; enter the estuary from the marsh when water levels drop  
254 below 5 cm) (Chick, Ruetz & Trexler 2004; Rehage & Loftus 2007; Trexler & Goss 2009;  
255 Parkos, Ruetz & Trexler 2011).

256

#### 257 Discrimination difference between blood and plasma

258 Studies quantifying isotopic discrimination values in sharks are limited. Hussey et al.  
259 (2010) investigated discrimination in captive sand tiger (*Carcharias taurus*; n = 3) and lemon  
260 sharks (*Negaprion brevirostris*; n = 1), however the only tissue they analyzed that could  
261 collected without lethal sampling was muscle tissue, which has a long isotopic half-life (~98  
262 days, MacNeil, Drouillard & Fisk 2006) and was not useful for our study. Both Kim et al.  
263 (2012) and Malpica-Cruz et al. (2012) investigated isotopic discrimination in captive leopard  
264 sharks (*Triakis semifasciata*; n = 6, n = 16, respectively). Kim et al. (2012) calculated  $\Delta^{13}\text{C}$   
265 plasma-blood values between 0.5-0.9‰. Despite sampling a variety of tissues, including blood,  
266 muscle, and fin tissue, Malpica-Cruz et al. (2012) did not collect plasma, and therefore data from  
267 this study was not useful for our analyses.

268 Tissue-specific incorporation of stable isotopes can be affected by variability in  
269 environmental conditions, and can vary between species (reviewed by Vander Zanden &  
270 Rasmussen 2001; Crawford, McDonald & Bearhop 2008; Newsome, Clementz & Koch 2010).  
271 Therefore, we used linear regression to estimate the inter-tissue discrimination difference  
272 between blood and plasma ( $\Delta^{13}\text{C}$  plasma-blood) in juvenile bull sharks, and compared this to  
273 values calculated by Kim et al. (2012). To do so, we plotted paired blood and plasma  $\delta^{13}\text{C}$   
274 values from 90 juvenile bull sharks against one another, and performed linear regression to

275 quantify the relationship between  $\delta^{13}\text{C}$  values (see Matich, Heithaus & Layman 2010 for further  
276 details). To test whether differences between  $\delta^{13}\text{C}$  values of blood and plasma varied across the  
277 observed range of  $\delta^{13}\text{C}$  values, we also used a t-test to determine if the slope of the best-fit-line  
278 was different from 1:1. We would expect deviation from a slope of 1:1 if  $\delta^{13}\text{C}$  discrimination  
279 varied with  $\delta^{13}\text{C}$  values of shark diets.

280

### 281 Temporal change in diet

282 In addition to providing an estimate of  $\Delta^{13}\text{C}$  plasma-blood, the regression plot of blood  
283 and plasma  $\delta^{13}\text{C}$  values described above also provided data that could be used to investigate  
284 dietary variability in sharks. Data points above the best fit line indicate plasma  $\delta^{13}\text{C}$  values more  
285 enriched (i.e. less negative) than predicted by the model, and data points below the regression  
286 line indicate plasma  $\delta^{13}\text{C}$  values more depleted (i.e. more negative) than predicted by the model  
287 (Fig. 3). In elasmobranchs, plasma has a faster isotopic turnover rate (half-life = 32 days, Kim et  
288 al. 2012) than blood (half-life = 61 days, MacNeil, Drouillard & Fisk 2006), and therefore  
289 dietary changes made by bull sharks should be detected by plasma isotope values considerably  
290 faster than blood isotope values. As such, plasma  $\delta^{13}\text{C}$  values more than 0.4‰ enriched above  
291  $\delta^{13}\text{C}$  values of blood (calculated  $\Delta^{13}\text{C}$  plasma-blood - See Results) indicate a recent change to  
292 either more marine food web-based diets or a change from feeding on marsh taxa to resident  
293 estuarine taxa. Plasma values more depleted than 0.4‰ above blood indicate either a recent  
294 change to more estuarine food web-based diets or a change from feeding on resident estuarine  
295 taxa to migratory marsh taxa (Fig. 2). Thus, we plotted the residuals from the regression analysis  
296 of blood and plasma  $\delta^{13}\text{C}$  against shark capture date [day of year (DOY)] to investigate temporal  
297 variability in the diets of bull sharks and to elucidate their potential use of the freshwater prey

308 pulse during the dry season. We then evaluated the effectiveness of using a piecewise function  
309 to describe the data against using the best fit line from linear or non-linear least squares  
310 regression (Toms & Lesperance 2003). We selected breakpoints at which  $\delta^{13}\text{C}$  residuals  
311 exhibited a notable change (DOY 128, 150, 163, 175, 213) and quantified the coefficient of  
312 determination for each model. We selected the piecewise model with the highest coefficient of  
313 determination and compared it to the coefficient of determination for linear and non-linear least  
314 squares fits to determine if it was significantly higher in order to choose the best overall model  
315 (Fisher 1921). To gain insight into general patterns of bull shark dietary changes in response to  
316 the prey pulse, we used ANOVA to test the effect of season on  $\delta^{13}\text{C}$  residual values. Post hoc  
317 Tukey's tests were used to test for significant differences across seasons.

318         Despite having isotope data from 2008-2012, we only used data from sharks caught from  
319 Oct 2008 to Dec 2009 because an extreme weather event in Jan 2010 significantly reduced the  
320 number of juvenile bull sharks in the estuary and changed the population structure (Matich &  
321 Heithaus 2012). It also possibly affected the community composition, and thereby trophic  
322 dynamics, of the ecosystem (Rehage et al. 2010). Additionally, we did not have acoustic  
323 tracking data for sharks after 2009 due to this extreme weather event, and therefore could not  
324 investigate the correlations between marsh water levels, shark movements, and stable isotope  
325 values. Future studies investigating interannual variation in shark trophic interactions and  
326 movements, however, would provide additional insights.

327         Due to the slow isotopic turnover rates of tissues in elasmobranchs (MacNeil, Drouillard  
328 & Fisk 2006; Logan & Lutcavage 2010; Kim et al. 2012) and the potential for maternal diets to  
329 be reflected in the tissues of newborns sharks (McMeans, Olin & Benz 2009; Matich, Heithaus  
330 & Layman 2010; Vaudo, Matich & Heithaus 2010), isotope values of bull sharks may not be



321 indicative of their current diet for individuals less than 90 days old (Belicka et al. 2012).  
322 Because bull sharks in the Shark River Estuary are likely born at 60-70 cm TL (based on  
323 captures of neonate individuals; Heithaus et al. 2009; Matich & Heithaus 2012) between May  
324 and August (based on the presence of umbilical scars; Curtis, Adams & Burgess 2011), and grow  
325 10-20 cm/year (based on recaptured individuals; Neer, Thompson & Carlson 2005), we only  
326 included tissues from individuals that were greater than 84 cm total length (at least one year of  
327 age) and individuals less than 85 cm TL that were caught between December and April with  
328 closed umbilical scars (at least 90 days old).

329         To determine if bull sharks changed their diets during the freshwater pulse, we developed  
330 a series of theoretical models to predict the differences in plasma and blood  $\delta^{13}\text{C}$  values to  
331 determine if plasma had recently become more enriched or depleted in  $\delta^{13}\text{C}$  in response to a  
332 dietary change (Fig. 4). We modeled six plausible dietary shifts. These included 1) estuarine  
333 prey  $\rightarrow$  marsh prey (E $\rightarrow$ F); 2) marine + estuarine prey  $\rightarrow$  estuarine + marsh prey (M+E $\rightarrow$ E+F);  
334 3) marine + estuarine prey  $\rightarrow$  marsh prey (M+E $\rightarrow$ F); 4) marine prey  $\rightarrow$  estuarine prey (M $\rightarrow$ E);  
335 5) marine prey  $\rightarrow$  estuarine + marsh prey (M $\rightarrow$  E+F); and 6) marine prey  $\rightarrow$  marsh prey (M $\rightarrow$ F)  
336 (Fig. 4b; Table 1). We used turnover data from MacNeil, Drouillard & Fisk (2006) (blood half-  
337 life = 61 days) and Kim et al. (2012) (plasma half-life = 32 days) to predict the rate of change in  
338 blood and plasma isotopes based on the differences in  $\delta^{13}\text{C}$  values of prey items from different  
339 food webs (mean  $\delta^{13}\text{C} \pm \text{SE}$ ; marine =  $-14.1 \pm 0.2\text{‰}$ , estuarine =  $-28.0 \pm 0.5\text{‰}$ , and marsh = -  
340  $30.5 \pm 0.5\text{‰}$ ). The models assumed that sharks would change their diets in response to marsh  
341 prey entering the system, with the timing of the modeled change based on the movements of bull  
342 sharks (see Results). Thus, the model assumed that shark diets changed when marsh water levels  
343 were  $\leq 0$  cm, which corresponds to the time when sharks significantly increased their use of the

344 upstream region [31 Mar 2009 (DOY 90)]. A second diet switch, to a diet similar to that before  
345 the prey pulse, was modeled to occur when water levels rose above 0 cm [3 Jun 2009 (DOY  
346 154)]. During predicted periods of dietary equilibrium (wet season), we assumed that blood and  
347 plasma values would differ by our calculated  $\Delta^{13}\text{C}$  plasma-blood (0.4‰; see Results).

348 Differences in  $\delta^{15}\text{N}$  values of marine, estuarine, and marsh taxa bull sharks may have fed  
349 upon (mean  $\delta^{13}\text{C} \pm \text{SE}$ ; marine =  $8.8 \pm 0.5\text{‰}$ , estuarine =  $9.0 \pm 0.5\text{‰}$ , and marsh =  $9.1 \pm 0.3\text{‰}$ )  
350 did not provide the same resolution as differences in  $\delta^{13}\text{C}$  values for taxa from each food web.  
351 Therefore, we only modeled changes in  $\delta^{13}\text{C}$ , rather than both  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$ . Stable isotope  
352 mixing models have become a popular analytical tool to investigate the trophic interactions of  
353 animals (reviewed by Layman et al. 2012), but mixing models do not provide adequate output to  
354 investigate temporal variability in the diets of individuals without repeated sampling, which is  
355 often difficult for highly mobile species, such as sharks. Our modeling approach, however,  
356 enabled us to quantify variability in the diets of each sampled shark in response to the freshwater  
357 prey pulse, and therefore we chose not to employ a mixing model.

358 Despite recent lab studies quantifying the turnover rates and discrimination values of  
359 blood and plasma stable isotopes in elasmobranchs (MacNeil, Drouillard & Fisk 2006; Logan &  
360 Lutcavage 2010; Kim et al. 2012; Malpica-Cruz et al. 2012), these processes can vary among  
361 similar species (reviewed by Vander Zanden & Rasmussen 2001; Crawford, McDonald &  
362 Bearhop 2008; Newsome, Clementz & Koch 2010). To investigate whether our estimates of  
363 discrimination and turnover rates might affect the performance of our models, we tested  
364 additional models in which we varied blood and plasma isotopic half-lives and  $\Delta^{13}\text{C}$  plasma-  
365 blood. We created models with the half-lives of blood and plasma decreased to half of published  
366 values (31 and 16 days, respectively) and increased to twice published values (122 days and 64

367 days, respectively; MacNeil, Drouillard & Fisk 2006; Kim et al. 2012) (Fig. 5a). We also  
368 created models with  $\Delta^{13}\text{C}$  plasma-blood of 0.9‰, 0.7‰, and 0.2‰, representing a range of  $\Delta^{13}\text{C}$   
369 plasma-blood values across the calculated values of Kim et al. (2012) (Fig. 5b). As such, we  
370 created six different models (each diet change scenario; see above) for 12 different treatments of  
371 isotopic half-life and  $\Delta^{13}\text{C}$  plasma-blood.

372 Because a piecewise function best described  $\delta^{13}\text{C}$  residuals across time (see Results), we  
373 used piecewise linear regression with the same breakpoint as the true  $\delta^{13}\text{C}$  residuals and DOY  
374 model (DOY = 169) to investigate the relationship between the predicted difference in  $\delta^{13}\text{C}$   
375 values (from theoretical models) and DOY. Because regression plots of predicted and true  
376 differences in  $\delta^{13}\text{C}$  values produced best fit lines with the same correlation coefficients and f-  
377 values for each diet change model across each isotopic half-life and  $\Delta^{13}\text{C}$  plasma-blood, we  
378 could not use traditional model selection. We therefore compared the best fit lines of the  
379 theoretical models to that of the model for true  $\delta^{13}\text{C}$  residuals and DOY. This approach allowed  
380 us to qualitatively select the best model(s) describing if and how bull sharks changed their  
381 trophic interactions in response to the prey pulse, and how isotopic half-life and  $\Delta^{13}\text{C}$  plasma-  
382 blood affected model selection. Criteria for qualitatively selecting the best theoretical models  
383 included 1) slopes of the piecewise functions with the same direction (positive or negative) as the  
384 model for true  $\delta^{13}\text{C}$  residuals and DOY; 2) slopes not significantly different from that of true  
385  $\delta^{13}\text{C}$  residuals and DOY (t-test); and 3) piecewise functions with the closest mean distance to the  
386 true  $\delta^{13}\text{C}$  residuals and DOY regression lines. ANOVA was used to test the effects of model,  
387 isotopic half-life, and  $\Delta^{13}\text{C}$  plasma-blood on mean distance from the true  $\delta^{13}\text{C}$  residuals and  
388 DOY piecewise function for theoretical models that passed the first two criteria. Post-hoc

389 Tukey's tests were used to test for significant differences across these factors. All statistical  
390 analyses were conducted in JMP 6.0.0.

391

## 392 **Results**

393 From 2008 to 2012, we captured 90 juvenile bull sharks. Twenty-three individuals  
394 between 71-142 cm total length (mean TL  $\pm$  SD = 102  $\pm$  22 cm) had acoustic transmitters  
395 surgically implanted in them, and were tracked from 10 Oct 2008 to 30 Nov 2009 for a total of  
396 5343 tracking days. Three individuals were not present for > 3 months within the system, and  
397 therefore were not included in movement analyses. Shark detections by upstream receivers  
398 varied by month, and were highest in April and May (DOY 91-151;  $R^2 = 0.59$ ,  $p < 0.01$ ; Fig. 6).  
399 During the dry season, there was a negative correlation between shark use of the upstream region  
400 and marsh water levels (DOY 60-148;  $R^2 = 0.52$ ,  $p < 0.01$ ; Fig 6), and the proportion of sharks  
401 detected in the upstream region was significantly higher when water levels were  $\leq 0$  cm (mean  $\pm$   
402 SD = 0.38  $\pm$  0.14) compared to > 0-5 (0.12  $\pm$  0.07) and > 0-10 cm (0.10  $\pm$  0.06) ( $t = 6.09$ ,  $p <$   
403 0.01;  $t = 8.54$ ,  $p < 0.01$ , respectively; Fig. 6).

404 Thirty-nine of the sampled sharks ( $n = 17$  males, 22 females) captured from Oct 2008 to  
405 Dec 2009 and ranging in size from 75-182 cm TL (mean TL  $\pm$  SD = 116.5  $\pm$  28.3 cm) were used  
406 to investigate seasonal shifts in shark diets relative to the 2009 pulse of prey from the marsh.  
407 Sharks had blood and plasma  $\delta^{13}\text{C}$  values that ranged from -17.5‰ to -26.5‰ (mean  $\delta^{13}\text{C} \pm$  SD  
408 = -22.9  $\pm$  2.4‰) and -17.8‰ to -25.3‰ (mean  $\delta^{13}\text{C} \pm$  SD = -22.4  $\pm$  2.3‰), respectively, and  
409 blood and plasma  $\delta^{15}\text{N}$  values that ranged from 10.5‰ to 12.8‰ (mean  $\delta^{15}\text{N} \pm$  SD = 11.6  $\pm$   
410 0.5‰) and 9.9‰ to 12.4‰ (mean  $\delta^{15}\text{N} \pm$  SD = 11.5  $\pm$  0.6‰), respectively (Fig. 2).

411 There was a significant, positive relationship between blood and plasma  $\delta^{13}\text{C}$  values with  
412 a high coefficient of determination (Fig. 3;  $R^2 = 0.81$ ,  $p < 0.01$ ), and the slope of the best fit line  
413 was not different from one (slope = 0.84,  $t_{(90)} = 0.18$ ). This suggests the mean difference  
414 between blood and plasma  $\delta^{13}\text{C}$  values (plasma was 0.4‰ greater than blood) was consistent  
415 across the  $\delta^{13}\text{C}$  range of the sampled sharks.

416 The  $\delta^{13}\text{C}$  residuals of bull sharks caught in 2008-2009 varied significantly with DOY and  
417 capture season. A piecewise function with a breakpoint at DOY 169 was significantly better  
418 than a polynomial fit or linear fit for the relationship between DOY and  $\delta^{13}\text{C}$  residuals (z-score =  
419 5.48,  $p < 0.01$ ; Fig. 7), with the magnitude of the slope for the first section (DOY 0-169) more  
420 than three times greater than the magnitude of the second section (DOY 169-365). Mean  $\delta^{13}\text{C}$   
421 residuals decreased significantly between the early dry season and the late dry season, and then  
422 increased from the late dry season to the early wet season (Fig. 8).

423 Changing the parameters of the models (isotopic half-life and  $\Delta^{13}\text{C}$  plasma-blood)  
424 changed their predictions of  $\delta^{13}\text{C}$  residuals. As the duration of isotopic half-life increased (i.e.  
425 from 0.5 half-lives to 2 half-lives), models predicted an increase in the duration of time  $\delta^{13}\text{C}$   
426 residuals were in a state of change in response to diet shifts, and as  $\Delta^{13}\text{C}$  plasma-blood increased  
427 (i.e from 0.2-0.9‰), models predicted greater positive  $\delta^{13}\text{C}$  residuals during non-pulse periods  
428 and smaller negative  $\delta^{13}\text{C}$  residuals during the prey pulse (Fig. 5). ANOVA revealed that model  
429 ( $F = 10.26$ ,  $p < 0.01$ ) and  $\Delta^{13}\text{C}$  plasma-blood ( $F = 14.08$ ,  $p < 0.01$ ) were significant factors  
430 explaining variability in mean distance between piecewise functions of theoretical models and  
431 the model of true  $\delta^{13}\text{C}$  residuals and DOY (Appendix 1). Models with  $\Delta^{13}\text{C}$  plasma-blood =  
432 0.7‰ and 0.9‰ had significantly lower mean distances from the true  $\delta^{13}\text{C}$  residuals and DOY  
433 piecewise function than discrimination differences of 0.2‰ and 0.4‰ (Fig. 9a), and models with

434  $\Delta^{13}\text{C}$  plasma-blood = 0.9‰ were the best fit (mean distance  $\pm$  SE =  $0.5 \pm 0.1\%$ ,  $0.4 \pm 0.1\%$  for  
435  $\Delta^{13}\text{C}$  plasma-blood = 0.7‰ and 0.9‰, respectively). Models 2 and 3 (marine + estuarine prey  
436  $\rightarrow$  estuarine + freshwater prey, and marine + estuarine prey  $\rightarrow$  freshwater prey, respectively) had  
437 the lowest mean distances from the  $\delta^{13}\text{C}$  residuals and DOY piecewise function (Fig. 9b), with  
438 model 2 having the best fit (distance  $\pm$  SE =  $0.4 \pm 0.1\%$ ,  $0.5 \pm 0.1\%$  for model 2 and model 3,  
439 respectively). This suggests that model 2 at  $\Delta^{13}\text{C}$  plasma-blood = 0.9‰ was the best model for  
440 predicting changes in  $\delta^{13}\text{C}$  values of blood and plasma, (Table 2, Appendix 1). Parameter  
441 estimates for isotopic half-life did not affect model performance ( $F = 0.90$ ,  $p = 0.42$ ).

442

## 443 **Discussion**

444 Seasonal resource pulses are important components of annual energy budgets for many  
445 species (reviewed by Otsfeld & Keesing 2000, Yang et al. 2008). For example, brown bears  
446 (*Ursus arctos*) in North American Pacific riparian ecosystems rely on predictable annual pulses  
447 of spawning salmon to sustain their biomass levels for overwinter hibernation (Naimen et al.  
448 2002, Helfield & Naimen 2006). Along the South African coastline, sardines (*Sardinops sagax*)  
449 make annual migrations into nearshore areas and serve as an important pulse of food for marine  
450 mammals, birds, bony fishes, and elasmobranchs (Dudley & Cliff 2010; O'Donoghue, Drapeau  
451 & Peddemors 2010). Within the Shark River Estuary, the influx of marsh taxa into upstream  
452 channels comprises a considerable proportion of the annual energy budget of teleost predators in  
453 the ecotone region (e.g. *Amia calva*, *Centropomus undecimalis*, *Micropterus salmoides*; Rehage  
454 & Loftus 2007; Boucek & Rehage *in press*), suggesting this resource pulse is likely a seasonally  
455 important component of estuarine food webs within the ecosystem. Our study shows that  
456 numerous juvenile bull sharks move upstream to take advantage of this influx of marsh prey,

457 despite relatively high levels of individual specialization within the population found in our  
458 previous work (Matich, Heithaus & Layman 2011).

459         Previously, we found that juvenile bull sharks in the Shark River Estuary show  
460 considerable inter-individual variation in trophic interactions, and many individuals (*ca.* 57%)  
461 showed relatively high degrees of specialization on one type of resource pool (i.e. marine food  
462 webs vs freshwater/estuarine food webs; Matich, Heithaus & Layman 2011). Our results from  
463 this study suggest the trophic interactions of some sharks in the estuary (i.e. those identified  
464 previously as specialists) are flexible, at least during the dry season when marsh taxa enter the  
465 system and provide an additional food source. Blood and plasma  $\delta^{13}\text{C}$  values (mean  $\pm$  SD = -  
466  $22.9 \pm 2.4\text{‰}$  and  $-22.4 \pm 2.3\text{‰}$ , respectively) suggest that many bull sharks fed on marine and  
467 freshwater/estuarine prey throughout the year, and during the wet and early dry seasons, sharks  
468 had  $\delta^{13}\text{C}$  residuals (plasma-blood) similar to our predictions attributed to  $\Delta^{13}\text{C}$  plasma-blood (*ca.*  
469  $0.3\text{-}0.9\text{‰}$ ), suggesting they had relatively stable diets. Yet, sharks had significantly lower  $\delta^{13}\text{C}$   
470 residuals during the late dry season (mean  $\pm$  SE =  $-0.5 \pm 0.4\text{‰}$ ), and model selection predicted  
471 diet switches from marine and estuarine prey to estuarine and marsh prey during the marsh prey  
472 pulse with a relatively rapid return to the previous diet at the terminus of the prey pulse when  
473 marsh prey were depleted, suggesting bull sharks fed on this seasonal resource pulse from  
474 freshwater marshes despite many individuals specializing on other resources outside this time  
475 period (Matich, Heithaus & Layman 2011).

476         Individual specialization has been hypothesized as a means to reduce intraspecific  
477 competition, especially in ecosystems with limited resources (reviewed by Bolnick et al. 2003).  
478 The Shark River Estuary is an oligotrophic system, and limited food abundance may be a driver  
479 of individual specialization in juvenile bull sharks (Matich, Heithaus & Layman 2011), however

480 the additional suite of resources from the marsh during the prey pulse may relax intraspecific  
481 competition for food. Apparently similar to bull sharks in the Shark River Estuary, specialist  
482 bluegill sunfishes (*Lepomis macrochirus*) were more successful foragers than generalists, but  
483 individual specialists exhibited flexibility in their habitat use and switched foraging behaviors  
484 when preferred resources became depleted (Werner, Mittlebach & Hall 1981). When preferred  
485 prey were scarce, silver perch (*Bidyanus bidyanus*) in aquaculture ponds switched from  
486 specializing on *Daphnia* to specializing on calanoids and insects, suggesting individual  
487 specialization may be a flexible trait in some populations (Warburton, Retif & Hume 1998). If  
488 bull shark trophic specialization is driven by density dependent processes, then the influx of  
489 marsh taxa into the system may considerably increase the availability of food, and reduce the  
490 need for sharks to have specialized diets when the prey pulse enters the estuary (Svanback &  
491 Persson 2004).

492

#### 493 *Drivers of bull shark behavior*

494 Previous studies in the Everglades suggest that large marsh taxa (> 8 cm TL) vacate the  
495 marsh in search of deep water refugia early in the dry season when water levels drop below 10  
496 cm in reference to elevation, and small marsh taxa (< 8 cm TL) enter the estuary later when  
497 water levels drop below 5 cm (Rehage & Loftus 2007; Trexler & Goss 2009). Thus, we  
498 predicted bull sharks would use the upstream region of the estuary early in the dry season to take  
499 advantage of all marsh taxa entering the system. Yet, bull sharks began using upstream areas  
500 heavily later than we predicted, which may be due to several reasons. Because many sharks in  
501 the estuary are less than three years old (Heithaus et al. 2009; Matich & Heithaus 2012), a lack of  
502 foraging experience may hinder their ability to detect when marsh prey are available (e.g.



503 Werner and Giliam 1984). Interannual variation in timing and magnitude of the prey pulse due  
504 to variation in precipitation and freshwater flow (Boucek & Rehage *personal communication*),  
505 may further reduce the ability of bull sharks to detect the start of this event. Unfortunately, our  
506 data set will not currently allow us to test these hypotheses.

507         Prey preference may also play a role in the timing of the bull sharks' responses to the  
508 prey pulse (Lanszki & Sallai 2006; Hawlena & Perez-Mellado 2009). If bull sharks preferred to  
509 eat large mesopredators like bass or bowfin, we would have expected them to use the upstream  
510 region earlier than observed, and their diets and isotope values would have changed accordingly.  
511 Instead, bull sharks did not significantly increase their use of the upstream region of the estuary  
512 until marsh water levels dropped below 0 cm, when all aquatic taxa have vacated the marsh.  
513 Thus, bull sharks may wait until the overall abundance of marsh taxa of all sizes in the system is  
514 relatively high, or they may be targeting smaller prey that arrive in the estuary later. Shark  $\delta^{15}\text{N}$   
515 values suggest that bull sharks likely targeted smaller prey from the marsh. Plasma and blood  
516  $\delta^{15}\text{N}$  values of bull sharks caught during the dry season (mean  $\pm$  SE =  $11.8 \pm 0.1\text{‰}$  and  $11.9 \pm$   
517  $0.2\text{‰}$ , respectively) were comparable to  $\delta^{15}\text{N}$  values of muscle tissue of other large aquatic  
518 predators like snook (mean  $\pm$  SE =  $11.3 \pm 0.3\text{‰}$ ) and bass (mean  $\pm$  SE =  $10.93 \pm 0.14\text{‰}$ ) that are  
519 known to feed on small marsh taxa. Therefore, bull sharks likely compete for with these large  
520 mesopredators for small prey that decline in abundance as the dry season progresses rather than  
521 consuming them (Boucek & Rehage *in press*). Comparison of  $\delta^{15}\text{N}$  values must be made  
522 cautiously, however. For example, muscle tissue in elasmobranchs has a slow turnover rate  
523 (half-life = 98 days, MacNeil, Drouillard & Fisk 2006), and thus we may not expect to detect  
524 large seasonal changes in  $\delta^{15}\text{N}$  values. Additionally,  $\delta^{15}\text{N}$  turnover and discrimination rates may  
525 vary with diet quality, trophic pathway, metabolic activity, and body size (reviewed by Vander

526 Zanden & Rasmussen 2001; Martinez del Rio et al. 2009; Hussey et al. 2012). Future studies  
527 incorporating stomach content analysis and fatty acid analysis should help further elucidate the  
528 importance of resource pulses to bull shark diets as well as intraspecific variation in the use of  
529 these resources.

530

### 531 *Alternative explanations*

532         Alternative explanations are unlikely to account for observed temporal variation in  
533 habitat use and  $\delta^{13}\text{C}$  values of sharks within the estuary. For example, shifts in habitat use by  
534 sharks could be driven by upstream movements of preferred prey (e.g. Ford et al. 1998; Rolstad,  
535 Loken & Rolstad 2000). Yet, if sharks were feeding on the same prey year round,  $\delta^{13}\text{C}$  residuals  
536 would be expected to remain similar during the year or exhibit longer lag-times if the prey of  
537 sharks had moved upstream to feed on the marsh prey pulse (i.e. the time for preferred prey to  
538 integrate marsh prey into tissues which would then be integrated into shark tissues).

539         Increased use of the upstream area by bull sharks when marsh water levels were  $\leq 0$  cm  
540 may have been driven by changes in environmental conditions. Juvenile bull sharks in other  
541 estuaries modify their space use in accordance with changes in salinity (e.g. Heupel &  
542 Simpfendorfer 2008; Froeschke, Stunz & Wildhaber 2010). Thus, bull sharks may have  
543 increased their use of the upstream region of the estuary during the dry season when salinities in  
544 areas further downstream increased and became higher than sharks preferred. However, salinity  
545 remains relatively low in the upstream region year-round (Heithaus et al. 2009, Rosenblatt &  
546 Heithaus 2011) and bull sharks are found in all areas of the estuary in all seasons (Matich &  
547 Heithaus 2012), suggesting physical factors are unlikely to be driving the significant increase in  
548 the use of the upstream area when marsh water levels are  $\leq 0$  cm.

549           Alterations in metabolic processes in response to environmental change can cause  
550 variability in stable isotope values (Kelly 2000; McCutchan et al. 2003; Vanderklift & Ponsard  
551 2003). Although bull sharks experience seasonal changes in salinity that may lead to changes in  
552 stable isotope values of tissues (Heithaus et al. 2009; Rosenblatt & Heithaus 2011), daily and  
553 weekly changes in salinity within the estuary would be expected to buffer a detectable change in  
554 isotope values attributed to osmoregulatory processes. Additionally, changing the isotopic half-  
555 lives and discrimination differences of our theoretical models did not affect the performance of  
556 our models or model selection (models 2 and 3 were the best models for all permutations),  
557 suggesting changes in metabolic processes attributed to environmental variability are unlikely to  
558 have produced the trends in  $\delta^{13}\text{C}$  observed during our study. As such, our results do indeed  
559 suggest sharks changed their diets during the dry season in response to the prey pulse, which  
560 may be a seasonally important source of nutrients and energy as observed in other predators  
561 within the system (Boucek & Rehage *in press*).

562

### 563 *Conclusions*

564           Stable isotope analysis is an attractive tool for ecologists because it can provide a time-  
565 integrated view of trophic interactions (Bearhop et al. 2004). While stable isotopes often provide  
566 only coarse information with regard to prey identity (reviewed by Gannes, O'Brien & Martinez  
567 del Rio 1997; Post 2002), employing this tool with complimentary approaches can be used to  
568 elucidate patterns and drivers of variability in trophic interactions and make predictions about  
569 how future conditions may lead to changes in food webs. Here we used a combination of stable  
570 isotope analysis, acoustic telemetry, and predictive modeling to elucidate changes in bull shark  
571 behavior in response to a resource pulse of taxa from adjacent marshland. Isotope data suggest

572 sharks increased the proportion marsh prey in their diets during the dry season, and movement  
573 data show that bull sharks increased their use of areas adjacent to freshwater marshes during this  
574 time. Annual variability in hydrology and planned changes in Everglades water management,  
575 however, may affect the importance of marsh taxa in the diets of bull sharks during the dry  
576 season.

577         Restoration efforts are planned to increase freshwater flow through the Everglades  
578 (CERP 2006), which will likely affect marsh water elevations (Obeysekera et al. 2011) and, in  
579 turn, the onset and duration of this resource pulse and the total biomass entering the Shark River  
580 Estuary. Increased freshwater flow and higher water levels in the marsh may lead to increased  
581 productivity, but may diminish the magnitude of the resource pulses into estuarine creeks, which  
582 could have negative consequences for the bull shark population and/or affect their ecological role  
583 within the ecosystem. If marsh taxa are not available within the estuary during the dry season,  
584 prey availability within the upper areas of the estuary may decrease and force bull sharks to  
585 increase their use of marine resources. This potential dietary shift may lead to decreased survival  
586 of young sharks, because downstream areas where marine taxa are most abundant are the riskiest  
587 habitats for small sharks to forage in due to high predation risk from large sharks (Heithaus et al.  
588 2009; Matich, Heithaus & Layman 2011). However, this shift in behavior may lead to upstream  
589 nutrient transport if juvenile sharks forage in areas where marine taxa are prominent, but rest  
590 upstream where large sharks are rarely found (Rosenblatt & Heithaus 2011). Additionally, if  
591 marsh taxa are not available to bull sharks, levels of individual specialization may further  
592 increase in the face of increased intraspecific competition (Matich, Heithaus & Layman 2011).  
593 Understanding how environmental variability currently affects the magnitude and timing of this  
594 pulse and the effects it has on aquatic communities is important for making predictions about

595 how changes in freshwater flow may alter slough communities in the Everglades. Using  
596 modeling approaches such as the one developed for this study can advance our understanding of  
597 temporal variation in trophic interactions, and provide predictions about how changes in the  
598 environment should affect food webs. Further research investigating the importance of resource  
599 pulses and disturbance regimes on the trophic dynamics of systems should increase our  
600 understanding of how predicted environmental changes due to natural and anthropogenic drivers  
601 may cause important ecological changes, and affect the role of predators within their respective  
602 ecosystems.

603

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620

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848 **Tables**

849 Table 1: Predicted  $\delta^{13}\text{C}$  values (in ‰) of prey in bull shark diets during periods of dietary  
850 equilibrium (wet season) and during dietary change (attributed to the freshwater prey pulse) used  
851 to predict  $\delta^{13}\text{C}$  residuals for the theoretical models. M = marine prey, E = estuarine prey, F =  
852 marsh prey entering channels during marsh dry down.

Model	$\delta^{13}\text{C}$ at equilibrium	$\delta^{13}\text{C}$ at prey pulse
1 (E→F)	-28.0	-30.5
2 (M+E→E+F)	-21.0	-29.3
3 (M+E→F)	-21.0	-30.5
4 (M→E)	-14.1	-28.0
5 (M→E+F)	-14.1	-29.3
6 (M→F)	-14.1	-30.5

853 Table 2: Mean distances  $\pm$  SE (in ‰) between actual  $\delta^{13}\text{C}$  residuals of bull shark blood and  
 854 plasma isotope values, and those predicted by theoretical models for each tissue-specific  
 855 discrimination difference between plasma and blood ( $\Delta^{13}\text{C}$  plasma-blood with plasma more  
 856 enriched for each scenario) to determine the best fit model(s) for sharks' diet change during the  
 857 freshwater prey pulse. Smaller distances indicate a better fit. M = marine prey, E = estuarine  
 858 prey, F = marsh prey entering channels during marsh dry down.

Model	$\delta^{13}\text{C}$ Discrimination difference			
	0.2‰	0.4‰	0.7‰	0.9‰
1 (E→F)	0.7 $\pm$ <0.1	0.6 $\pm$ 0.1	0.5 $\pm$ 0.1	0.5 $\pm$ 0.1
2 (M+E→E+F)	0.7 $\pm$ <0.1	0.5 $\pm$ <0.1	0.2 $\pm$ <0.1	0.1 $\pm$ <0.1
3 (M+E→F)	0.8 $\pm$ <0.1	0.5 $\pm$ <0.1	0.3 $\pm$ <0.1	0.2 $\pm$ <0.1
4 (M→E)	1.0 $\pm$ 0.1	0.8 $\pm$ 0.1	0.6 $\pm$ 0.1	0.5 $\pm$ 0.2
5 (M→E+F)	1.0 $\pm$ 0.1	0.8 $\pm$ 0.1	0.7 $\pm$ 0.1	0.6 $\pm$ 0.2
6 (M→F)	1.1 $\pm$ 0.1	0.9 $\pm$ 0.1	0.8 $\pm$ 0.2	0.7 $\pm$ 0.2

859

860 **Figure captions:**

861 Figure 1: The study occurred in the Shark River Estuary, Florida, USA. The star represents  
862 United States Geological Survey water station 252820080505400, which was used to define  
863 seasons. Locations of acoustic receivers are indicated by white circles, and the white rectangle  
864 encompasses the upstream region where freshwater prey enter the estuary during marsh dry  
865 down.

866 Figure 2: Mean isotope values for producers and consumers in the Shark River Estuary and  
867 adjacent marine waters from Fry & Smith (2002), Chasar et al. (2005), Williams & Trexler  
868 (2006), and our own sampling. Producers and consumers from the freshwater/estuarine food  
869 web are black, those from the marine food web are gray, and migratory marsh taxa are white.  
870 Producers are pluses (+), primary consumers are triangles (▲), secondary consumers are squares  
871 (■), tertiary consumers are circles (●), and bull sharks (blood isotope values) are diamonds (◆).

872 Figure 3: Linear regression of paired blood and plasma  $\delta^{13}\text{C}$  values. The mean difference  
873 between blood and plasma  $\delta^{13}\text{C}$  values (0.4‰) serves as an approximation for the difference in  
874  $\delta^{13}\text{C}$  discrimination between blood and plasma in bull sharks. The location of data points relative  
875 to the regression line provides insights into whether an individual's diet has become more  
876 enriched in  $\delta^{13}\text{C}$  or more depleted in  $\delta^{13}\text{C}$  than predicted by differences in discrimination factors  
877 of blood and plasma. Open circles are data from sharks caught in 2008-2009, and closed  
878 diamonds are data from sharks caught in 2010-2012. Only sharks caught in 2008-2009 were  
879 considered for temporal analysis.

880 Figure 4: Model predictions for changes in a)  $\delta^{13}\text{C}$  values of plasma and blood and b)  $\delta^{13}\text{C}$   
881 residuals if bull sharks switched to using freshwater prey during the dry season. If bull shark  
882 diets consist of resident estuarine taxa and are at equilibrium during the wet season, mean blood  
883  $\delta^{13}\text{C}$  values should be -28.0‰ and mean plasma  $\delta^{13}\text{C}$  values should be -27.6‰. When marsh  
884 taxa enter the estuary during the dry season, if bull sharks switch to feeding on marsh taxa  
885 (Model 1), plasma  $\delta^{13}\text{C}$  values will become more depleted faster than blood  $\delta^{13}\text{C}$  values because  
886 plasma  $\delta^{13}\text{C}$  turnover (~32 day half-life) is faster in elasmobranchs than blood  $\delta^{13}\text{C}$  turnover (~  
887 61 day half-life). In this scenario, differences between plasma and blood  $\delta^{13}\text{C}$  values are  
888 predicted to switch from being positive to negative on DOY 112 and then revert to being positive  
889 on DOY 162 after marsh taxa have become depleted and bull sharks return to feeding on  
890 estuarine taxa. Note that in b) the inconsistencies at the ends of the diet switch periods (near  
891 DOY 148 and 200) are attributed to the different rates of change in plasma and blood  $\delta^{13}\text{C}$   
892 (plasma approaches its asymptote much sooner than blood).

893 Figure 5: Effects of changing a) isotopic half-life at  $\Delta^{13}\text{C}$  plasma-blood = 0.4‰ and b)  $\Delta^{13}\text{C}$   
894 plasma-blood at one half-life on predictions of diet-change model 2.

895

896 Figure 6: Relationship between marsh water elevation (gray line) at United States Geological  
897 Survey water station 252820080505400 and the proportion of sharks detected by upstream  
898 receivers per day (black line) from 10 Oct 2008 (DOY 284) to 31 Nov 2009 (DOY 335).

899 Figure 7: Temporal variation in  $\delta^{13}\text{C}$  residuals. The black lines are predicted residuals based on a  
900 piecewise function and the gray dashed line is the predicted change in the difference between  
901 plasma and blood  $\delta^{13}\text{C}$  values for model 2 (marine + estuarine prey  $\rightarrow$  estuarine + freshwater  
902 prey attributed) at one isotopic half-life and  $\Delta^{13}\text{C}$  plasma-blood = 0.9‰, which was the best  
903 model for predicting changes in  $\delta^{13}\text{C}$  residuals. Model selection was not influenced by  
904 assumptions about isotopic half life or  $\Delta^{13}\text{C}$ .

905 Figure 8: Seasonal variation in mean  $\delta^{13}\text{C}$  residuals. Error bars are  $\pm$  SE, and bars with different  
906 letters are significantly different based on post hoc Tukey's tests.

907 Figure 9: Mean differences between actual  $\delta^{13}\text{C}$  residuals and those predicted by a) all models at  
908 each  $\delta^{13}\text{C}$  discrimination difference between tissues and b) across each model for all  
909 discrimination differences and half-lives. Data are means and error bars are  $\pm$  SE, and bars with  
910 different letters are significantly different based on post hoc Tukey's tests.

Figure 1:

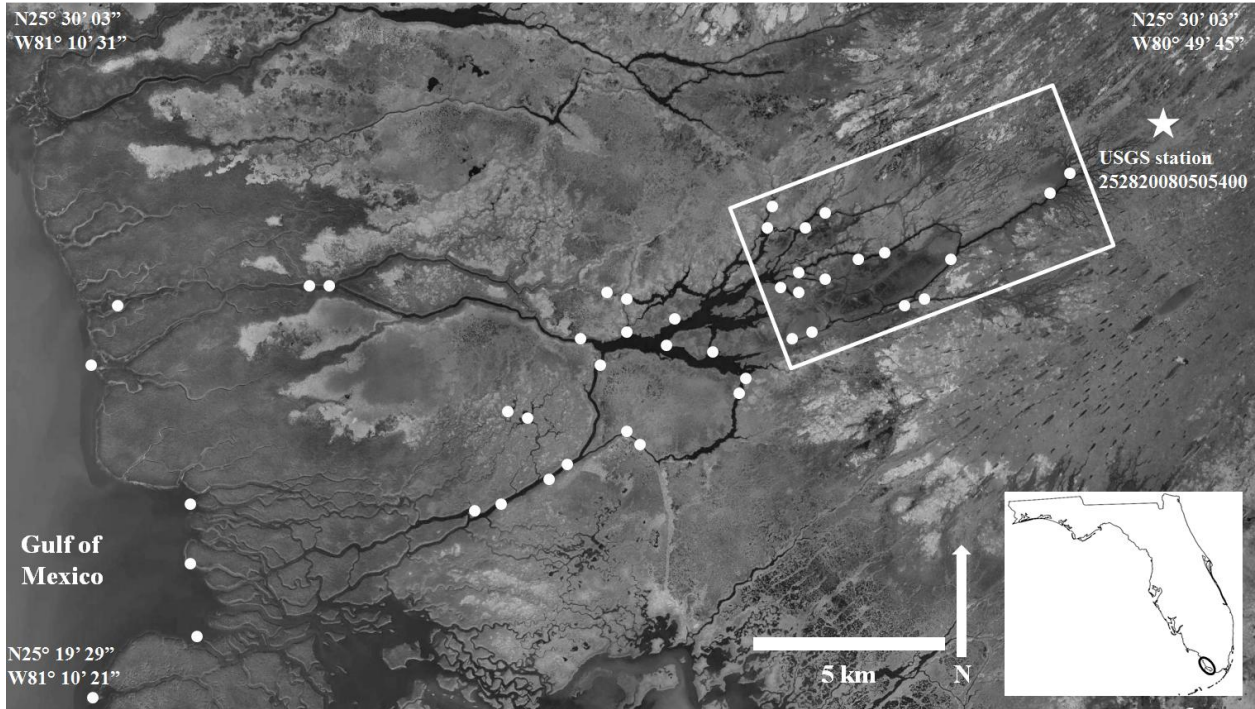


Figure 2:

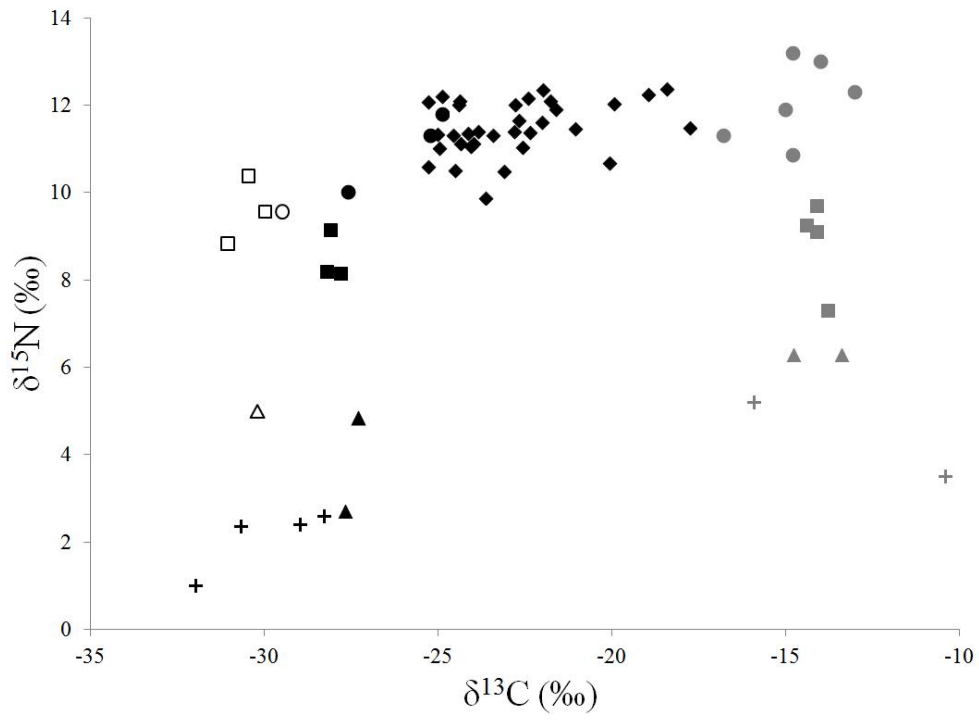


Figure 3:

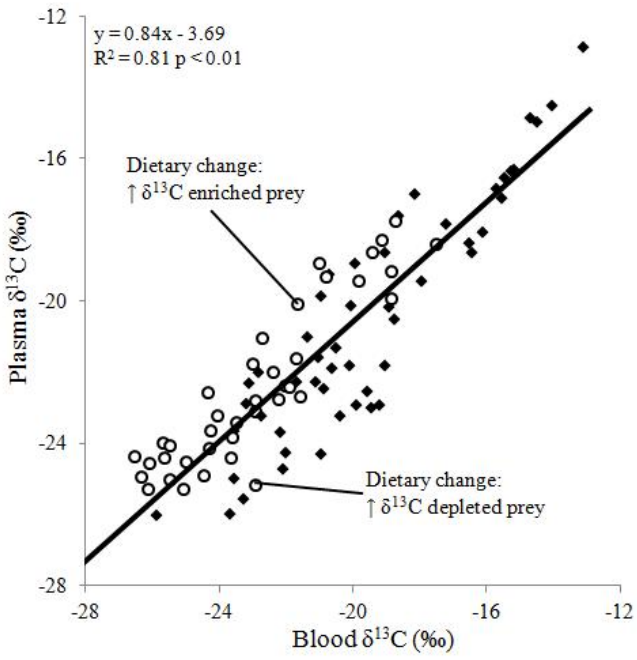


Figure 4:

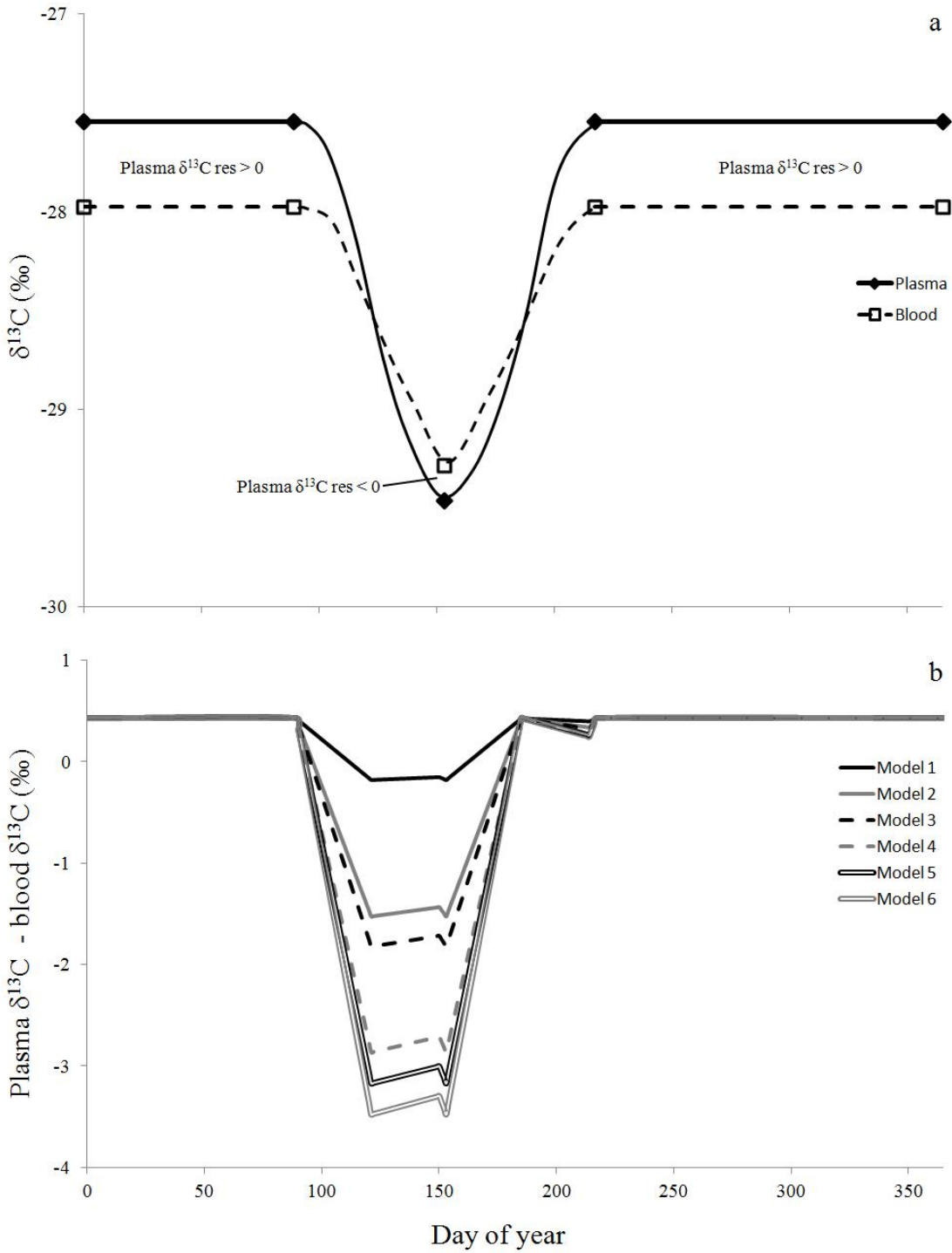




Figure 5:

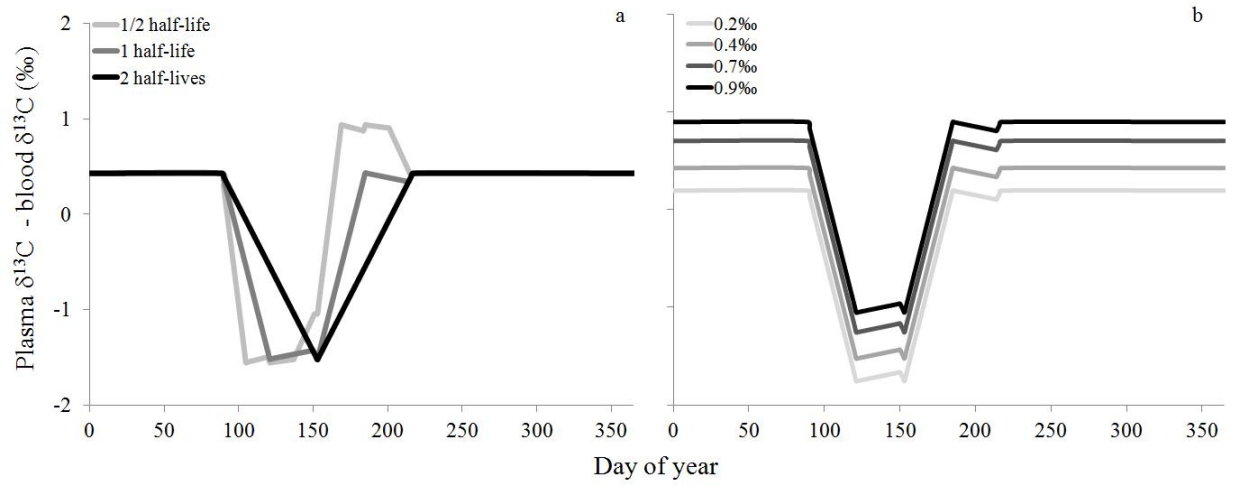


Figure 6:

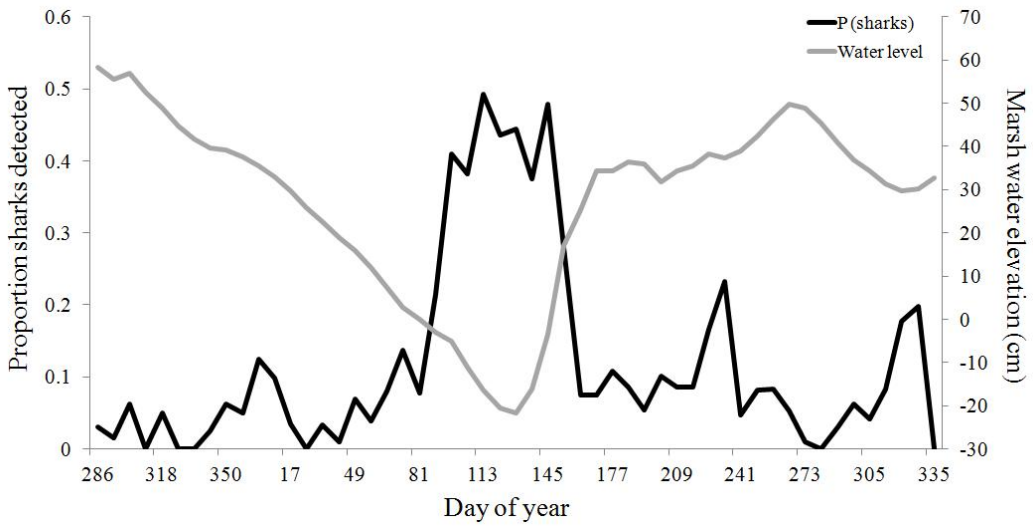


Figure 7:

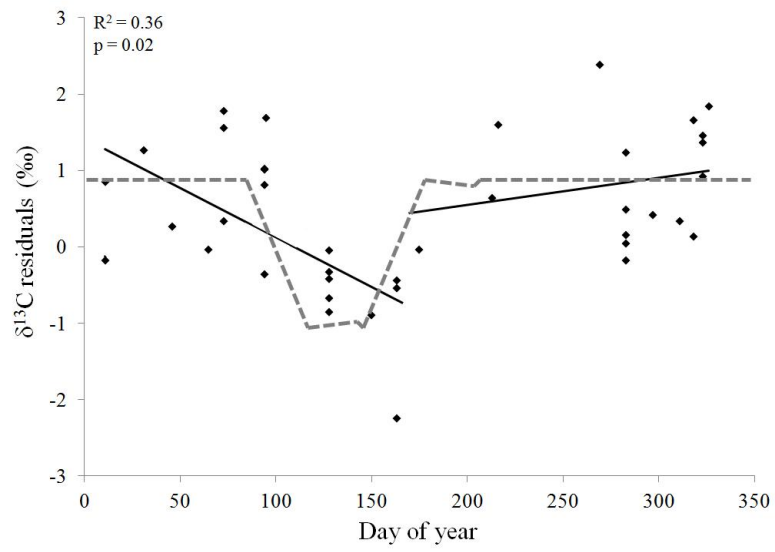


Figure 8:

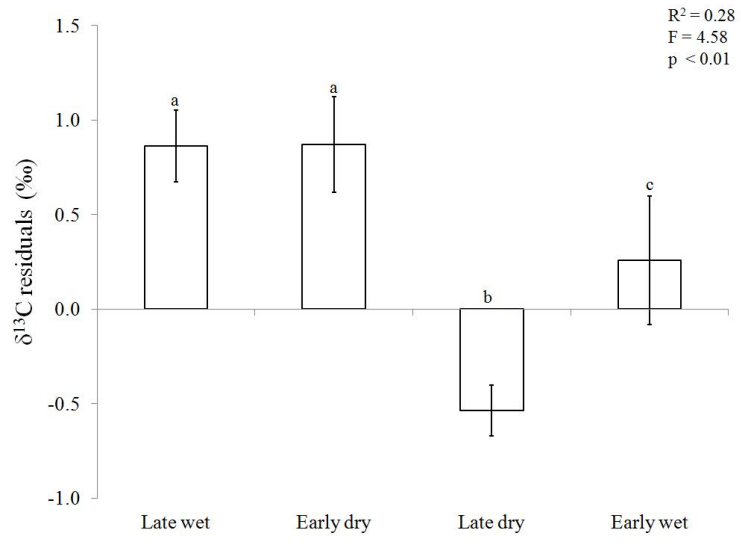


Figure 9:

