



17 Marine protected areas (MPAs) have been recommended as an essential conservation tool for  
18 ecosystems and fisheries, with MPA coverage increasing four-fold globally since 2000. Despite  
19 the increased usage of MPAs, empirical results and scientists are divided on the effectiveness of  
20 MPAs to reach their conservation and management goals. In 2009, the South Atlantic Fishery  
21 Management Council established eight deep-water partially-protected MPAs off the southeast  
22 United States (SEUS) Atlantic coast with the goal of protecting long-lived, deep-water, reef-  
23 associated fishery species. Data collected during fish trapping from 2000 to 2018 as part of a  
24 fisheries-independent survey were used to evaluate the impacts of protection on a subset of fish  
25 species in three of these MPAs. There was a modest positive shift to larger fish observed in two  
26 of the MPAs and to older fish for one species, *Pagrus pagrus* (Linnaeus, 1758), in one of the  
27 MPAs relative to adjacent fished areas; however, there was either no change or a decrease in  
28 managed reef fish abundance in each MPA relative to adjacent fished areas. Based on these  
29 metrics, it does not appear that the SEUS MPAs have yet been effective at protecting managed  
30 reef fish species. Given that these MPAs have low enforcement, future assessments should  
31 examine compliance within the SEUS MPAs to determine if lack of success is due to illegal  
32 fishing, species examined, or MPA design before making a final determination if deep-water  
33 MPAs are an effective strategy for fisheries managers in the SEUS.

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35 **Keywords:** Marine Protected Areas; Partially-Protected; Deep-Water; Reef Fish; Southeast  
36 United States; Atlantic Ocean

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## 41 **1. Introduction**

42 Marine protected areas (MPAs) have become widely used by fisheries managers to  
43 increase abundance, biomass, biodiversity, and body sizes of previously targeted fish inside  
44 MPAs by limiting fishing mortality; in some cases, this has promoted spillover into adjacent  
45 regions (Russ & Alcala 1996; Babcock *et al.* 1999; Pomeroy *et al.* 2004; Russ, Stockwell &  
46 Alcala 2005; Di Lorenzo *et al.* 2016). Angulo-Valdés & Hatcher (2010) proposed many other  
47 potential benefits of MPAs that encompass a broad scope of ecological, social, and economic  
48 values. Because of these potential benefits, in 2016 the International Union for Conservation of  
49 Nature pushed to increase the establishment of MPAs, aiming to conserve at least 30% of the  
50 global ocean by 2030 (<https://portals.iucn.org/library/node/46467>).

51 In 2009, the South Atlantic Fishery Management Council (SAFMC) established eight  
52 deep-water MPAs along the continental shelf break off the southeast United States (SEUS)  
53 Atlantic coast. The establishment of the eight MPAs was a nearly two-decade long process, with  
54 the final placement and design of the MPAs representing a compromise between fishers and the  
55 SAFMC to meet fisheries management goals while minimizing economic and social impacts  
56 (SAFMC 2006). The intent of the designations of the SEUS MPAs was to protect a portion of  
57 long-lived, deep-water, reef-associated fishery species by prohibiting bottom fishing inside the  
58 MPAs. Many species of reef fish often share habitat preferences, and as a result it is common to  
59 catch and subsequently discard one species when another species is targeted. Release mortality in  
60 reef fish species is higher in deeper water due to increased barotrauma (Wilson & Burns 1996;  
61 Burns & Restrepo 2002; Burns *et al.* 2002; Rudershausen *et al.* 2007; Runde and Buckel 2018);  
62 thus, the establishment of MPAs in deep-water was pursued. As partially-protected MPAs,

63 fishing for (bottom fishing), possessing, or retaining reef fish species is prohibited, but trolling  
64 and harvesting of other species, such as mackerel, marlin, or tuna, is allowed (SAFMC 2016).  
65 The MPAs also have a transit provision allowing for fishers with reef fish species onboard their  
66 vessels to traverse the MPA if their fishing gear is stowed (SAFMC 2016). It was believed that  
67 by solely prohibiting bottom fishing, these MPAs would, in turn, decrease catch and bycatch  
68 mortality and protect spawning areas for reef fish, ultimately promoting spillover into fishable  
69 areas (SAFMC 2016). Additionally, the placement of the SEUS MPAs along the shelf could  
70 provide potential sources of fish larvae as they are transported throughout the SEUS via the Gulf  
71 Stream and inner and mid shelf currents if enhanced spawning occurs in these MPAs (Hare &  
72 Walsh 2007; Leshner 2008; Govoni *et al.* 2013; SAFMC 2016).

73           Since the establishment of the SEUS MPAs, only one study has been published that  
74 examined the efficacy of these MPAs (Bacheler *et al.* 2016). Using underwater videos from  
75 manned and unmanned submersibles (2001-2014 time series), Bacheler *et al.* (2016) determined  
76 that there was no increase in the number of reef fish species or density of some managed reef fish  
77 species in SEUS MPAs regionally, and in two SEUS MPAs individually, six years after their  
78 implementation. At the time of that study, the MPAs were only six years old and thus changes to  
79 the fish community may have not had enough time to accrue. A meta-analysis of MPAs globally  
80 showed that MPA age (>10 years) was one of five key criteria influencing MPA effectiveness  
81 (Edgar *et al.* 2014). In addition, previous studies have shown that the recovery time for long-  
82 lived reef fish abundance and biomass within MPAs can be on the scale of decades (McClanahan  
83 & Mangi 2000; Russ *et al.* 2005; Russ & Alcala 2010). This suggests that continued evaluation  
84 of the SEUS MPAs is needed as their lifespans increase.

85 Another reason Bachelier *et al.* (2016) observed no MPA effect may be due to the  
86 biological indicators used to evaluate SEUS MPA effectiveness (species richness and individual  
87 density). Indicators used to evaluate protection effectiveness vary across MPAs and will depend  
88 on MPA goals and objectives (Pomeroy *et al.* 2004; Pelletier *et al.* 2005; Fox *et al.* 2014;  
89 Gallacher *et al.* 2016). One of the primary goals of the SEUS MPAs, as stated by the SAFMC, is  
90 to protect the size, age, and genetic structure of deep-water species susceptible to overfishing  
91 (SAFMC 2016). The Pelletier *et al.* (2005) review of MPA metrics showed that density and body  
92 size distributions were the most effective indicators of rehabilitation of population and age  
93 structure in fish. In addition, ecological and environmental benefits following MPA  
94 establishment occur over different timeframes, with increased size, age, abundance, and biomass  
95 expected shortly after MPA establishment with biodiversity and community composition benefits  
96 expected later (Ward *et al.* 2001; Stelzenmuller & Pinnegar 2011). As such, demographic (size  
97 and age) indicators, in addition to abundance, should be used to evaluate the effectiveness of the  
98 SEUS MPAs.

99 We evaluated the impact of protection in three of the eight SEUS MPAs using  
100 abundance, size, and age of managed reef fish species. We used trap catch data from fishery-  
101 independent surveys from the year 2000 (nine years pre-establishment) to 2018 (nine years post-  
102 establishment) from within the MPAs and adjacent fished areas. We focused on the commonly  
103 encountered managed reef fish species to provide an overall view of protection impacts and a  
104 more specific focus on Red Porgy, *Pagrus pagrus* (Linnaeus, 1758), a relatively short-lived but  
105 common reef fish species that is likely to benefit from protection due to its overfished status  
106 (SEDAR 2020).

107

108 **2. Methods**

109           This study used data collected by the Marine Resources Monitoring Assessment and  
110 Prediction (MARMAP) program from 2000 to 2008 and the Southeast Reef Fish Survey  
111 (SERFS) from 2010 to 2018 to examine changes in abundance, size, and age of managed reef  
112 fish species in response to the implementation of three deep-water MPAs off the southeast  
113 United States (SEUS) Atlantic coast. Initially, regional reef fish monitoring in the SEUS was  
114 conducted solely by the MARMAP program. MARMAP has since been joined by two other  
115 programs, the Southeast Area Monitoring and Assessment Program – South Atlantic (SEAMAP-  
116 SA) and the Southeast Fishery Independent Survey (SEFIS). This collaboration is now referred  
117 to as SERFS, with each collaborator using gear and methodologies consistent with historical  
118 MARMAP sampling (Bacheler *et al.* 2014; Smart *et al.* 2015). The SERFS sampling covers the  
119 continental shelf and shelf edge between Cape Hatteras, North Carolina, and St. Lucie Inlet,  
120 Florida. The survey is the main contributor of fishery-independent data to SEUS reef fish species  
121 stock assessments, collecting information on relative abundance, distribution and demographics  
122 for managed (and some unmanaged) reef fish species. Although this survey was not designed to  
123 explicitly test MPA effectiveness, its coverage of known reef habitat sites inside and outside the  
124 MPAs as well as sampling pre- and post-MPA establishment makes it one of few data sets  
125 available to assess reef fish response to SEUS MPA establishment. In addition, to the best of our  
126 knowledge, it is the only data set available that has monitored changes in abundance, size, and  
127 age of reef fish, important metrics to assess the effectiveness of the SEUS MPAs.

128 **2.1 Study Sites**

129           The size of the MPAs established by the SAFMC in 2009 range from 30 to 515 km<sup>2</sup>, with  
130 water depths from 50 m to 300 m, and boundaries encompassing a variety of outer shelf, shelf-  
131 edge, and upper slope habitats. MARMAP and SERFS samples include five of the nine deep-

132 water MPAs. Due to sampling characteristics such as the geographic scope of the survey,  
133 available funding, and survey priorities, three of the five MPAs sampled most consistently were  
134 analyzed in this study: Northern South Carolina (NSC), Edisto (ED), and North Florida (NF)  
135 (Figure 1; Table 1). Samples collected from 2000 to 2018, excluding 2009 (year of MPA  
136 implementation), were used for inside-outside and before-after comparison for the NSC and  
137 Edisto MPAs. Due to low sampling effort prior to the establishment of the NF MPA, only  
138 samples collected from 2010 to 2018 were used, and only for an inside-outside comparison.

139 Sites used for inside-outside MPA comparisons were selected from adjacent fished areas  
140 (<55 km from MPA boundary), with sufficient sample sizes and similar depth and distance to the  
141 continental shelf edge. Depth and distance to shelf edge were considered in the designation of  
142 adjacent fished areas as they have been shown to be dominating factors structuring fish  
143 assemblage in the SEUS (Glasgow 2017). Additionally, proximity to shelf edge ensured that  
144 MPAs and adjacent fished sites were approximately equidistant to the coastline and fishing ports.  
145 These adjacent fished comparison areas covered a larger spatial extent than their respective MPA  
146 comparison areas, as sampling outside the MPA was not as concentrated.

## 147 **2.2 Data Collection**

148 As part of SERFS standard sampling, chevron traps (see Collins 1990) were used to  
149 sample along the SEUS continental shelf in depths ranging from 9 to 115 m. Each year, a subset  
150 of stations was selected randomly from known reef stations identified for monitoring in a manner  
151 such that no station selected in a given year was closer than 200 m to any other selected station.  
152 Prior to deployment each chevron trap was baited with a combination of 40 whole or cut  
153 clupeids, with *Brevoortia* spp. most often used. Generally, chevron traps were deployed in sets of  
154 six and soaked for approximately 90 min, during which time a CTD cast was done. Gear  
155 deployments occurred during daylight hours (no earlier than 30 minutes after sunrise and

156 retrieved no later than 30 minutes before sunset). Depth (m), latitude, longitude, bottom  
157 temperature (C), time of day, date, and soak time were recorded as part of the standard SERFS  
158 sampling protocol for each trap deployment.

159         Beginning in 2011, all chevron traps used by SERFS were equipped with a camera  
160 mounted above and facing away from the trap opening (trap mouth) and a camera mounted  
161 above and facing away from the back of the traps (trap back) to record habitat characteristics.  
162 All camera settings remained constant throughout a given sampling year and were set to record  
163 continuous video during the entire soak period of the chevron traps until memory cards filled or  
164 batteries ran out of power. Following gear retrieval, videos were uploaded to external hard  
165 drives, processed on shore, and used in this study for characterizing habitat type. Habitat  
166 characterization followed SERFS standard protocols (Bacheler *et al.* 2014). Habitat was coded  
167 from each video immediately upon the trap stabilizing on the sea floor, noting dominant surface  
168 geologic and biotic components: substrate density, size, and relief, and dominant biota height,  
169 type, and density. Prior to this study, habitat coding had been done by trained SEFIS personnel  
170 for 2011-2017 and was used for this study. Samples collected in 2018 were coded for habitat by  
171 trained MARMAP/SEAMAP-SA personnel following the same protocol.

172         Once a set of traps was fully retrieved, all fish caught in each trap underwent a Length  
173 Frequency work-up (LF). LF occurred shortly after retrieval and consisted of identifying and  
174 counting all fish in each trap to species level or the lowest possible taxon, then weighing all fish  
175 of a given species per trap as an aggregate (biomass in g wet weight) and measuring each  
176 individual fish. Abundance was calculated for each species per trap based on the number  
177 measured. Prior to 2012, lengths were measured as either total length or fork length in cm and  
178 since 2012 all lengths for all species were measured as total length in cm. Fork lengths were



179 converted to total length for this study based on conversions developed by Bublely *et al.* (2019).  
180 During LF, specimens of priority species were retained for additional processing to collect  
181 biological samples (Life History or LH processing). Any fish not retained for the LH processing  
182 were degassed (vented) as necessary and released.

183 On-board, as part of LH processing, both sagittal otoliths were removed from Red Porgy,  
184 *Pagrus pagrus* (Linnaeus, 1758), the only species examined in this study for changes in age  
185 structure. Both sagittal otoliths were placed in dry coin envelopes until further processing on  
186 shore. The left sagitta (or right if the left was broken or unavailable) was used in age estimation.  
187 Otolith sectioning followed standard MARMAP protocols (Smart *et al.* 2015). Otolith  
188 increments were determined by the number of alternating translucent and opaque bands and were  
189 counted independently by two trained readers without knowledge of specimen length, mass,  
190 location, date of capture, or results of the other reader. Upon completion of independent reads,  
191 increment counts were compared and in cases where readers disagreed, readers simultaneously  
192 viewed the sections and attempted to reach a consensus. If consensus could not be reached, those  
193 specimens were eliminated from analyses.

### 194 **2.3 Study Design**

195 This study used a *post hoc* sampling design to control for differences in habitat structural  
196 complexity and uneven sample sizes between MPAs and their adjacent fished comparison areas.  
197 In many cases, more stations were available for analysis in the adjacent fished areas than in the  
198 MPA, leading to uneven sample sizes and a risk of comparing sites of dissimilar habitat  
199 complexity. Since the implementation of camera-mounted chevron traps in 2011, it was clear  
200 that among chevron stations in the SERFS sampling universe, there was considerable variation in  
201 reef habitat structural complexity. Camera data were not consistently available for habitat  
202 characterization for trap deployments before 2011, however chevron sites sampled 2000-2010

203 consisted of the same stations used for 2011-2018 sampling. Previous work done by MARMAP,  
204 using a series of repeated measures analyses of variance, showed that there were no significant  
205 changes in some habitat characteristics (relief, substrate density, and biota density) over time for  
206 traps deployed at the same station from 2011 to 2017 (Smart *et al.* 2020). These results supported  
207 *post hoc* assignment of habitat scores for samples collected 2000-2010 in MPA and adjacent  
208 fished areas based on station number. This was done by averaging substrate and biota density  
209 and taking the mode of vertical relief for station numbers that were sampled multiple times ( $\geq 3$ )  
210 from 2011 to 2018. For stations that did not have habitat codes assigned, habitat classification  
211 from the nearest station with habitat characterized was used. Nearby stations generally have a  
212 high degree of correlation in habitat type for the three characteristics included here based on  
213 spatial autocorrelation results (Moran's I for biota density = 0.697, substrate density = 0.717, and  
214 vertical relief = 0.709 and all p-values <0.001) and because the survey specifically targets hard  
215 bottom habitats. In addition, the average distance between stations with significantly different  
216 habitat types was >120 m while distance to the nearest station used as a proxy for habitat was on  
217 average 51 m.

218         Average station depth along with the three proxies for habitat complexity described  
219 above (vertical relief, substrate density, and biota density) were used to select adjacent fished  
220 area deployments with similar habitat characteristics to MPA deployments for analyses. These  
221 categories were selected as they have been shown to be some of the dominant habitat features  
222 structuring reef fish assemblage in the SEUS (Glasgow 2017). To avoid subjectivity in selecting  
223 deployments, habitat similarity was determined by cluster analysis in R using the cluster package  
224 (Maechler *et al.* 2019). Because depth, substrate density, and biota density are continuous and

225 vertical relief is categorical, a Gower's Distance test was used to produce a distance metric for  
226 cluster analysis (Gower 1971; Borcard *et al.* 2011).

227 Gower's tests were done for each year of sampling and for each MPA and adjacent fished  
228 area pair as not every station was sampled in each year of the survey due to the random selection.  
229 Continuous variables (substrate and biota density) were square root transformed to meet  
230 normality assumptions. Partitioning around Medoids algorithm was then used to create clusters  
231 comprising MPA and adjacent fished area sites from each Gower's distance metric. The number  
232 of clusters in which to partition the distance metric was determined by the highest average  
233 silhouette width. Silhouette width is a measure of how similar an object (in this study the object  
234 is each deployment) is to its own cluster and how distinct it is from other clusters. This grouped  
235 deployments (MPA and fished) into clusters of low to high structural habitat complexity with  
236 similar depth. Deployments were then selected from the adjacent fished area based on their  
237 respective silhouette width, with deployments having the greatest silhouette width being  
238 selected. The number of adjacent fished deployments chosen from each cluster were based on the  
239 number of MPA deployments that occurred within that cluster to produce even sample sizes. If  
240 there were fewer adjacent fished deployments in a cluster than MPA deployments, then adjacent  
241 fished area deployments with the greatest silhouette width from the nearest neighbor cluster were  
242 selected. Essentially, this clustering approach provided a structured, reproducible, unbiased way  
243 of selecting comparative deployments outside of the MPAs and producing even samples sizes for  
244 MPAs and fished areas. The approach also ensured we compared sites of similar habitat  
245 complexity (and quality) by minimizing the possibility of comparing sites of high reef habitat  
246 complexity inside the MPA to sites of low habitat complexity in the adjacent fished area, or vice  
247 versa.

## 248 **2.4 Data Analysis**

249           The combined relative abundance of the ten most commonly caught managed reef fish  
250 species (RFS) in chevron traps for each MPA served as a proxy for the effects of MPAs on  
251 managed reef fish species in general, even though the exact composition of the RFS groups  
252 differed among MPAs (Table 2). Most common is defined in this paper as highest frequency of  
253 occurrence and abundance in catches. Changes in Red Porgy abundance in response to MPA  
254 implementation was chosen for species-specific analyses because they were commonly caught in  
255 all three MPAs. Also, due to Red Porgy's overfished status (SEDAR 2020), relatively short  
256 longevity (< 18 years), and early maturing life history patterns (~100% mature at age 4; Wyanski  
257 *et al.* 2019), Red Porgy was expected to respond more quickly to MPA implementation than  
258 other, longer lived SEUS managed reef fish species (Polacheck 1990; Hilborn *et al.* 2004;  
259 Rassweiler *et al.* 2014), such as the Speckled Hind (*Epinephelus drummondhayi*, Goode & Bean,  
260 1878), which may live up to 80 years and reach maturity at 7 years (Andrews *et al.* 2013; Kobara  
261 *et al.* 2017). Using a trap catch data set permitted us to examine changes in abundance, while  
262 also examining changes in size and age. Despite these benefits, traps, like any survey gear type,  
263 are not able to efficiently collect all the information desired to monitor changes in an area,  
264 because some species are too large or less willing to enter traps or too small to be retained  
265 (Bacheler *et al.* 2013a). Thus, community-based parameters to assess the effects of MPA  
266 implementation, such as richness, evenness, or changes in community structure, were not  
267 examined in this study.

268           We used multiple regression models to examine changes in abundance of RFS and Red  
269 Porgy in response to protection while factoring out relevant covariates. Specifically, we fit the  
270 following linear model:

$$\begin{aligned} \log(\text{abundance} + 1) = & \beta_0 + \beta_1(\text{region}) + \beta_2(\text{location}) + \beta_3(\text{time}) + \beta_4(\text{location} * \text{time}) + \\ & \beta_4(\text{temperature}) + \beta_5(\text{depth}) + \beta_6(\text{relief}) + \beta_7(\text{substrate density}) + \beta_8(\text{biota density}) + \text{offset}(\text{trap} \\ & \text{soak time}) + \epsilon ; \epsilon \sim \mathbf{N}(0, \sigma^2) \end{aligned}$$

271 The primary variables of interest in this global model were: *region* which specified the Northern  
 272 South Carolina, Edisto, or Northern Florida region, *location* which identified whether a given  
 273 deployment was inside or outside of the MPA, *time* which denoted whether the deployment was  
 274 before or after MPA implementation (year 2009), and the *location-by-time* interaction which was  
 275 necessary to identify if deployments located within the MPA were different after protection was  
 276 implemented. The other variables in the model were covariates that we wished to statistically  
 277 control for including: *temperature* the bottom temperature (°C) from the paired CTD cast, *depth*  
 278 the sampling depth (m) of the deployment, and *relief*, *substrate density*, and *biota density* were  
 279 the average habitat characteristics for the station number where the deployment occurred. We  
 280 included *trap soak time* as an offset term inside the model to account for variability in the  
 281 sampling effort as some deviation from the targeted 90-minute soak time occurred which could  
 282 affect abundance within the traps (Zuur *et al.* 2009; Bachelier *et al.* 2013b). An offset was used  
 283 rather than dividing the abundance by soak time to allow variability to be captured in the model  
 284 and allow us to investigate a wider range of error distributions appropriate for count data. Since  
 285 geographic location (latitude and longitude) is specific to each MPA and adjacent fished  
 286 comparison area, we did not include that as a predictor variable. We also fit this model to each  
 287 region separately but dropped the region explanatory variable.

291 We compared the fit (i.e., negative log likelihood) of the Gaussian linear model on the  
 292  $\log(x + 1)$  transformed abundance to General Linear Models with Poisson, negative binomial,  
 293 and zero-inflated error structures. The negative binomial and Gaussian models consistently

294 outperformed the other models. Diagnostic plots for assessing model assumptions (normality and  
295 variance) and model prediction, showed that Gaussian structure was the most appropriate with  
296 one fewer fitted parameter. We used Analysis of Variance (ANOVAs; Quinn and Keough 2002)  
297 (type 3) to test for significance of explanatory variables and overall model fit at the  $\alpha = 0.05$   
298 level.

299         It is possible that there may be a delayed effect in protection on abundance, which  
300 prompted an examination of trends of abundance over narrower time intervals. General linear  
301 power analyses (GLP) performed on RFS and Red Porgy abundance suggested that a sample size  
302 of 38 to 160 was needed to test for significance. These results prompted years to be grouped into  
303 three-year periods (2000-2002, 2003-2005, 2006-2008, 2010-2012, 2013-2015, 2016-2018) to  
304 ensure an adequate sample size for examining trends over time.

305         Changes in size were examined for RFS and Red Porgy. RFS for each MPA comparison  
306 were grouped into four periods to limit pair-wise comparisons: 2000-2004, 2005-2008, 2010-  
307 2014, and 2015-2018. Due to limited sample size prior to MPA establishment, Red Porgy total  
308 lengths were grouped for each MPA comparison into three periods: 2000-2008, 2010-2014, and  
309 2015-2018. Changes in age were analyzed in response to MPA establishment for Red Porgy  
310 from 2000 to 2017 and were grouped into three periods based on sample size: 2000-2008, 2010-  
311 2013, and 2014-2017. To examine changes in Red Porgy age distributions, we compared changes  
312 in the percent of Red Porgy expected to be fully recruited to the fisheries as these would indicate  
313 a benefit to fisheries and a change in age structure. Red Porgy age at full recruitment to chevron  
314 trap ( $>$  age 3) was determined using length-age data collected from 2000 to 2017 and resulting  
315 catch curve and was consistent with Hood & Johnson (2000). To maintain an even sample size  
316 and habitat characteristics, the same deployments used in the abundance analyses for each of the

317 MPAs were used for size and age comparisons. Before-after and inside-outside changes in size  
318 distributions and percent of fully recruited fish were assessed for significance using a series of  
319 one-sided t-tests (assuming unequal variance) and chi-square tests, respectively. Because NF  
320 MPA samples were limited prior to MPA establishment, analyses of change in size and percent  
321 recruited were only examined post implementation.

### 322 **3. Results**

#### 323 **3.1 Reef Fish Species Abundance**

324 Generally, the Reef Fish Species (RFS) linear models, both globally and for each MPA,  
325 were significant ( $p < 0.05$ ), but had low  $R^2$  values ( $< 0.13$ ; Table 3). None of the models we  
326 examined were able to capture the degree of dispersion observed in fish abundance  
327 (Supplemental Figure 1). Across all three regions, the global models showed that RFS abundance  
328 was significantly ( $p < 0.05$ ) lower inside the MPAs than outside and that there was no significant  
329 ( $p > 0.05$ ) interaction effect (Figure 2; Table 3). In the North South Carolina (NSC) area,  
330 abundance was significantly ( $p < 0.05$ ) greater outside the MPA, pre- and post-MPA  
331 implementation, and exhibited a negative trend in abundance inside the MPA post establishment  
332 (Figure 3a). In the Edisto (ED) area, there was a significant ( $p < 0.05$ ) positive interaction effect,  
333 with greater abundance inside the MPA post-establishment relative to the adjacent fished area  
334 and pre-MPA establishment levels. From an examination of periods post-MPA implementation,  
335 it is apparent that the positive interaction was driven by a large increase in abundance in 2013-  
336 2015 (Figure 3b). In the most recent period of sampling, ED MPA abundance level was not  
337 significantly different ( $p > 0.05$ ) from abundance levels inside the MPA for any period pre-  
338 establishment or for 2010-2012. In the North Florida (NF) area, abundance inside the MPA was

339 significantly lower ( $p < 0.05$ ) than the adjacent fished area after MPA establishment (Figure 2;  
340 Figure 3c).

### 341 **3.2 Red Porgy Abundance**

342 Generally, the Red Porgy linear models, both globally and for each MPA, were  
343 significant ( $p < 0.05$ ), but had low  $R^2$  values ( $< 0.13$ ; Table 3). Global models showed that Red  
344 Porgy abundance was significantly ( $p < 0.05$ ) greater inside than outside the MPAs post  
345 establishment, but there was no significant ( $p > 0.05$ ) interaction effect (Figure 2; Table 3). In the  
346 NSC area, there was no significant inside-outside, before-after or interaction effects. Red Porgy  
347 abundance for periods post-establishment show overlapping standard error bars and no  
348 significant ( $p > 0.05$ ) differences between the NSC MPA and adjacent fished area (Figure 4a). In  
349 the ED area, the ED MPA had significantly greater Red Porgy abundance than the adjacent  
350 fished area, both before and after MPA implementation, but showed no significant interaction  
351 effect (Figure 3; Figure 4b). In the NF area, there was no significant ( $P > 0.05$ ) difference in Red  
352 Porgy abundance inside or outside the MPA post establishment (Figure 3; Figure 4c).

### 353 **3.3 Size-Distributions of RFS and Red Porgy**

354 RFS mean total length (TL) inside the NSC MPA area was similar to the adjacent area for  
355 the period immediately prior to MPA implementation ( $p = 1.0$ ) and in the most recent period post  
356 MPA implementation ( $p = 0.99$ ; Figure 5a). Mean RFS TL was significantly greater inside the  
357 ED MPA compared to the adjacent fished area for all periods ( $p < 0.001$ ) except 2005-2008 ( $p =$   
358  $1.0$ , Figure 5b). In 2000-2004, RFS mean TL inside the ED MPA was 1.8 cm greater than the  
359 adjacent area. In 2015-2018, RFS mean TL inside the ED MPA was 4.7 cm greater than the  
360 adjacent fished site, suggesting an increase in the size difference over time. For both periods post  
361 NF MPA establishment, mean TL was significantly greater ( $p < 0.001$ ) inside the MPA than the



362 adjacent fished area, with difference in mean TL increasing over time from 2.8 cm to 7.1 cm  
363 (Figure 5c).

364 Red Porgy mean TL was similar inside the NSC MPA and in the adjacent area, regardless  
365 of the period ( $p > 0.59$ , Figure 6a). Red Porgy mean TL was significantly greater inside the ED  
366 MPA compared to the ED adjacent fished area for all periods (2000-2008,  $p = 0.05$ ; 2010-2014  
367 & 2014-2018,  $p < 0.001$ , Figure 6b). In both periods post MPA establishment, RFS mean TL  
368 inside the ED MPA was ~3 cm greater than the adjacent fished area. Finally, Red Porgy mean  
369 TL was significantly greater inside the NF MPA compared to the adjacent fished area by 4.1 cm  
370 during the final period of sampling ( $p < 0.001$ ; Figure 6c).

### 371 **3.4 Age Distributions of Red Porgy**

372 Overall, percentage of fully recruited Red Porgy in 2014-2017 was significantly lower  
373 inside the NSC MPA, significantly greater inside the NF MPA, and not significantly different  
374 inside the ED MPA compared to their respective adjacent fished areas. Prior to NSC MPA  
375 establishment, percentage of fully recruited Red Porgy was significantly greater inside the NSC  
376 MPA than outside ( $p < 0.001$ ; Figure 7a). For the period immediately following NSC MPA  
377 establishment, there was no significant difference in percentage of fully recruited Red Porgy  
378 inside or outside the MPA. During the most recent period (2014-2017), percent of fully recruited  
379 Red Porgy was significantly greater outside the NSC MPA than inside (67.7% and 50.0%  
380 respectively,  $p < 0.01$ ), a reversal of the pre-establishment pattern. Prior to ED MPA  
381 establishment, there was no significant difference in percentage of fully recruited Red Porgy  
382 inside and outside the ED MPA (Figure 7b). For the period immediately following ED MPA  
383 establishment, percentage of fully recruited Red Porgy was greater inside the MPA than in the  
384 adjacent fished area (45.7% and 18.6%, respectively;  $p=0.002$ ). During the most recent period

385 (2014-2017) both areas experienced an increase to ~80% fully recruited with no difference  
386 between the protected and adjacent fished area. For the period immediately following NF MPA  
387 establishment, percentage of fully recruited Red Porgy was lower inside the MPA than outside  
388 (46.8% and 68%, respectively;  $p < 0.001$ ; Figure 7c). The most recent period (2014-2017),  
389 percentage of fully recruited Red Porgy was greater inside the NF MPA than the adjacent fished  
390 area (75.6% and 61.5%, respectively;  $p = 0.02$ ), a reversal of the immediate post-establishment  
391 pattern.

#### 392 **4. Discussion**

393 Our results suggest that the southeast United States (SEUS) MPAs examined here have  
394 not had a positive effect on exploited reef fish species, with a few exceptions. We did not  
395 observe fish abundances (Reef Fish Species (RFS) or Red Porgy) increasing inside of the MPAs  
396 after protection was implemented. Linear models showed no change, or in some cases appeared  
397 to exhibit a negative trend, in abundance of RFS and Red Porgy, both globally and for each of  
398 the three MPAs and adjacent fished area comparisons. The only exception to this was the Edisto  
399 MPA, which displayed a significant interaction effect, with RFS increasing inside the MPA over  
400 time relative to the adjacent fished area. However, upon further examination, trends in RFS  
401 abundances post-establishment showed that in the most recent sampling period there was no  
402 significant difference in RFS abundance inside and outside the MPA or before and after MPA  
403 implementation. The extraordinarily high catches in the ED MPA from 2013-2015 most likely  
404 drove the ED MPA positive interaction effect and the nearly significant ( $p \sim 0.07$ ) interaction term  
405 for the global model. These results suggest that as of 2018, the SEUS MPAs have not (yet) been  
406 (or no longer are) effective in increasing abundances of managed reef fish species.  
407 Unfortunately, this is not uncommon among MPAs, with many partially-protected MPAs such as

408 these being deemed ineffective and underperforming relative to no-take MPAs (Burke *et al.*  
409 2011; Denny & Babcock 2004; Lester & Halpern 2008; Edgar *et al.* 2014).

410 Further examination of some biological indicators did show a positive response to MPA  
411 implementation. A significant positive before-after and inside-outside response was observed for  
412 the size of RFS and Red Porgy in ED MPA. A significant positive inside-outside response was  
413 also observed for the size of RFS and Red Porgy and age of Red Porgy in the NF MPA. While  
414 fish length comparisons between protected and adjacent fished areas may only differ by a few  
415 centimeters, this could be the difference between a legal- and sublegal-sized fish. Some of the  
416 more commonly caught reef fish in the SEUS, Vermilion Snapper (*Rhomboplites aurorubens*,  
417 Cuvier, 1829), Gray Triggerfish (*Balistes capriscus*, Gmelin, 1789), Black Sea Bass  
418 (*Centropristis striata*, Linnaeus, 1758), and Red Porgy have TL minimum limits between 30.5  
419 and 35.5 cm, which falls within the range of lengths observed in this study. When TLs of Red  
420 Porgy were examined (TL minimum limit = 35.5 cm), ED and NF MPAs both showed a  
421 significantly higher proportion (72% & 88%, respectively) of TLs over the minimum size limit  
422 post MPA establishment compared to adjacent fished area (62% & 64.5%, respectively; Figure  
423 7). Additionally, the proportion of Red Porgy over the minimum size limit for ED MPA  
424 increased from 58% just prior to MPA establishment to 72% for the most recent period post  
425 MPA establishment. From a management perspective, this could be an important result  
426 increasing the availability of legally-sized fish in the region if spillover occurs. These size  
427 increases within the NF and ED MPA support previous findings on both the benefits of protected  
428 areas and the moderate size gains exhibited in partially protected MPAs relative to no-take areas  
429 (Babcock *et al.* 1999; Unsworth *et al.* 2007; Malcolm *et al.* 2015; Harasti *et al.* 2018).

430 Three of the reef fish species used in the RFS analysis had changes to their minimum size  
431 limit between 2000 and 2018: Black Sea Bass, Grey Triggerfish, and Red Snapper, (*Lutjanus*  
432 *campechanus*, Poey, 1860). The Black Sea Bass size limit increased by 2.54 cm in 2012, while a  
433 minimum size limit of 30.5 cm was imposed for Gray Triggerfish in 2016 and the size limit for  
434 Red Snapper was removed in 2011. While we do not expect these modest changes in size limit to  
435 affect our results, it is possible that it could provide a bias for the before-after comparisons,  
436 where observations are driven by size limit changes rather than the implementation of the MPA.  
437 However, since the data used in this study was collected as part of a fisheries-independent  
438 survey, which is not subjected to size restrictions, inside-outside comparisons for the RFS  
439 analysis should be able to identify changes in size related to MPA establishment rather than  
440 fishing mortality.

441 One possible reason why no positive effect was observed for abundance was the short  
442 span of time since the MPAs were implemented (Russ *et al.* 2005; Russ & Alcala 2010; Edgar *et al.*  
443 *al.* 2014). Age of MPA typically is recognized as one of the most important criteria influencing  
444 MPA effectiveness (Claudet *et al.* 2008; Molloy *et al.* 2009; Edgar *et al.* 2014). Yet, studies  
445 differ on the minimum time needed to be effective, from only a few years (Halpern & Warner  
446 2002; Unsworth *et al.* 2007), to a decade (Edgar *et al.* 2014), to several decades (MacNeil *et al.*  
447 2015). This difference can be attributed to differing life-history strategies of protected fish  
448 species, with a greater delay in protection response for slow-growing and long-lived species  
449 (Molloy *et al.* 2009; Claudet *et al.* 2010; Starr *et al.* 2015). Catches here were generally  
450 dominated by faster maturing and shorter-lived reef fish species, such as Vermilion Snapper and  
451 Red Porgy (Potts *et al.* 1998; Hood & Johnson 2000). Based on the relatively early age of  
452 maturity and faster growth of most species encountered, the MPAs appear to have been

453 established long enough for an increase in abundance to be observed within the boundaries of  
454 MPA sampled, provided that there is indeed reduced fishing pressure and sufficient habitat in  
455 these areas.

456 Another major driver of MPA effectiveness is overall size (Claudet *et al.* 2008; Fenberg  
457 *et al.* 2012; Edgar *et al.* 2014). In order to be effective, MPAs must be large enough to ensure  
458 that fish are spending enough time inside the protected area to reduce fishing mortality and meet  
459 management goals. The three MPAs studied range in size from 170 to 340 km<sup>2</sup>. Unfortunately,  
460 few movement pattern studies have been conducted on deeper reef fish in the SEUS to better  
461 understand their home ranges because of the difficulty of tagging these fish. Studies of reef fish  
462 species in shallower regions have shown that daily home ranges are well within the size limits of  
463 these MPAs, with some grouper, such as Red Grouper (*Epinephelus morio*, Valenciennes, 1828),  
464 Black Grouper (*Mycteroperca bonaci*, Poey, 1860), and Gag (*Mycteroperca microlepis*, Goode  
465 and Bean, 1879) (Coleman *et al.* 2011; Farmer & Ault 2011), and snapper, such as Mutton  
466 Snapper (*Lutjanus analis*, Cuvier, 1828) and Yellowtail Snapper (*Ocyurus chrysurus*, Bloch,  
467 1791) (Farmer & Ault 2011), and Red Porgy (Afonso *et al.* 2009) exhibiting home ranges of less  
468 than 8 km<sup>2</sup>. However, some of the managed reef fish species undergo ontogenetic migrations,  
469 moving from coastal to offshore habitats, as well as migrating great distances to spawning  
470 aggregations potentially taking them well outside the protection of these MPAs (Ross & Moser  
471 1995; Coleman *et al.* 2000). Given the variability of movement patterns of studied managed reef  
472 fish species and general lack of information for deep reef fish species it is unclear what MPA  
473 size is needed to ensure that SEUS managed reef fish species remain inside the MPA to benefit  
474 from protection.

475 MPA placement and design could also have been a factor affecting MPA effectiveness.  
476 The SEUS MPAs were established as a result of a nearly two-decade long process and reflect a  
477 compromise between fishers and fisheries managers (SAFMC 2006). This compromise resulted  
478 in the MPAs being placed in locations to “lessen social and economic impacts”, which explains  
479 why they were designed perpendicular to the shelf contours rather than parallel (SAFMC 2006),  
480 which would have encompassed larger areas of reef habitat. This effectively limits the  
481 availability of on-shelf and shelf edge reef habitat in these SEUS MPAs and concentrated reef  
482 habitat to the western boundaries of the MPAs. This is particularly evident with the NSC MPA  
483 where shelf reef habitat is concentrated in the northwestern corner of the MPA, resulting in the  
484 smallest on-shelf sampling area of the three MPAs studied (NSC ~ 72 km<sup>2</sup>, ED ~ 170 km<sup>2</sup>, and  
485 NF ~ 110 km<sup>2</sup>). This limitation in sampling coverage and habitat may explain why the NSC  
486 MPA showed no positive effect for any metrics examined, and in some cases exhibited a  
487 negative response. Numerous studies have shown that the amount of available habitat and spatial  
488 distribution of habitat contribute significantly to MPA effectiveness, helping to maintain species  
489 diversity and provide shelter/refuge for species across the MPA (Foley et al. 2010; Gaines et al.  
490 2010; Heyns-Veale et al. 2019). Additionally, studies have shown that fishing pressure typically  
491 increases at the margins of MPAs as fishers expect a spillover of larger/older individuals or more  
492 fish moving out of a protected area, known as the edge effect or “fishing the line” (Carr *et al.*  
493 2003; McLeod *et al.* 2009; Green *et al.* 2014). This possible increased fishing pressure on the  
494 edge of the MPA, coupled with the fact that reef fish habitat is concentrated near the MPA  
495 boundary suggests that fish in the NSC MPA area may have been experiencing greater fishing  
496 pressure than the other two MPAs studied.

497           Based on age, size, and design of MPAs alone it is still unclear why no positive response  
498 in abundance was found in the North Florida and Edisto MPAs. This may be attributed partially  
499 to the lack of effective enforcement and compliance, a major factor affecting MPA success  
500 (Lundquist & Granek 2005; Guidetti *et al.* 2008; Edgar *et al.* 2014; Di Franco *et al.* 2016). In the  
501 most recent deep-water MPA management plan by the South Atlantic Fisheries Management  
502 Council (SAFMC), seven of the eight MPAs were given a rating of “low” for enforceability by  
503 the US Coast Guard (SAFMC 2016). This is driven by two factors: (1) distances from shore to  
504 MPAs and (2) the partial protection status of the MPAs allowing certain fishing activities to take  
505 place, making on-site enforcement, rather than flyovers, a necessity to determine if fishing is  
506 unlawful or not (SAFMC 2016). Since the MPAs have been established, not a single citation has  
507 been issued for illegal fishing (USCG, personal communication). This is not necessarily an  
508 indication of the level of infractions but may reflect the limited patrolling and difficulty  
509 associated with enforcing regulations. The consequences of instituting partially protected areas  
510 instead of full no-take reserves have been shown widely in other regions, with no take reserves  
511 outperforming partially protected areas (Francour 1994; Denny & Babcock 2004; Sciberras *et al.*  
512 2013; Edgar *et al.* 2014; Ballantine 2014). Because of the low enforceability of the SEUS MPAs,  
513 it is nearly impossible to assess compliance within the protected areas, which is a major problem  
514 moving forward with evaluations of SEUS MPAs as a management strategy in SEUS Atlantic  
515 region.

516           Sampling design could have influenced our results and explain why metrics did not  
517 support that SEUS MPAs provide effective protection for reef fish. Because our samples were  
518 collected as part of a regional survey, they may be limited relative to a study designed to  
519 specifically test MPA effectiveness. However, it is the only standardized sampling available

520 before and after MPA establishment that provides information on reef fish size and age  
521 compositions, in addition to abundance. Chevron trap sampling by MARMAP and SERFS was  
522 limited to previously established sites with known reef habitat <115 m depth. This restricted  
523 sampling to on-shelf, shallower regions of the MPAs relative to the full spatial scope of the  
524 MPAs. Reef fish species assemblages' change with depth in the SEUS (Glasgow 2017), and the  
525 results of this study therefore may not be as applicable to reef fish distributed in deeper, off-shelf  
526 portions of the MPAs. It should be noted that limited high-resolution mapping of the MPAs has  
527 shown much less reef habitat in the deeper regions of the MPAs than where sampling was  
528 conducted (SAFMC 2016), and the effects of protection provided to deeper reef fish may be  
529 inconsequential due to the limited availability of deep reef habitat. Additionally, reef fish  
530 distributed in off-shelf waters tend to be slower-growing and longer-lived (Coleman *et al.* 2000;  
531 Sabetian 2003; Costa *et al.* 2012; Bubley *et al.* 2019), thus one would expect these species, and  
532 subsequently deeper regions of the MPAs, to respond more slowly to protection (Molloy *et al.*  
533 2009; Starr *et al.* 2015). Here we focused on species expected to be more responsive due to their  
534 shorter life spans and times to maturity and still found limited evidence of MPA effectiveness.  
535 Lastly, although our samples were collected as part of a larger survey, our novel *post hoc*  
536 stratification approach did mitigate at least some uncertainty that results were driven by  
537 differences in habitat complexity or sampling effort.

538         The SEUS MPAs were implemented as a fisheries management tool. The desired goal  
539 was that by protecting suitable reef fish habitat in the MPAs, this would result in higher  
540 abundance and biomass of exploited reef fish and increase the number of individuals reaching  
541 reproductive maturity within the MPA, promoting the spillover of fish out of the protected  
542 boundaries into the surrounding areas. From size and age metrics, our study indicates that the



543 SEUS MPAs are protecting reef fish, permitting more fish to reach larger sizes and older ages  
544 which may relate to their reproductive output; however, none of the MPAs exhibited an increase  
545 in abundance and, in some cases, abundance showed a decline. We believe this may negate the  
546 modest gains in size and age observed and would not result in a significant MPA effect on  
547 reproductive potential or population recovery. In addition, patterns of effectiveness were not  
548 consistent across MPAs. For example, the NSC MPA showed no positive fish size responses to  
549 protection, while only the NF MPA showed increases in fish age relative to its adjacent fished  
550 area. Unfortunately, until compliance within MPAs has been assessed, it will be difficult to  
551 determine if deep-water MPAs, or the current deep-water MPA orientations and protection level,  
552 are an effective strategy for protecting reef fish over the long term. If fisheries managers in the  
553 SEUS Atlantic region wish to use MPAs as a management tool, it is important to continue to  
554 understand why an increase in abundance has not been observed and what is driving the  
555 differences among the MPAs responses post-establishment.

556

#### 557 **CRedit author statement**

558 **Chris Pickens:** Formal Analysis, Visualization, Investigation, Writing – Original Draft,  
559 Methodology, **Tracey Smart:** Supervision, Data Curation, Writing – Review and Editing,  
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575

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811 **Tables**

812 Table 1. Sampling distributions of three southeast United States (SEUS) marine protected areas  
813 (MPAs): Northern South Carolina (NSC), Edisto (ED), and North Florida (NF). Provided are the  
814 area and depths sampled as compared to the total area and depth range of the MPAs.

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<b>MPA</b>	<b>Sampling Area (km<sup>2</sup>)</b>	<b>Boundary Area (km<sup>2</sup>)</b>	<b>Sampling Depth Range (m)</b>	<b>Boundary Depth Range (m)</b>
NSC	72	170	45-68	45-170
ED	170	170	45-60	45-140
NF	110	340	45-64	45-380

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825 Table 2. Most commonly caught managed reef fish species in Northern South Carolina (NSC),  
 826 Edisto (ED), and North Florida (NF) marine protected areas and adjacent fished areas. x = ten  
 827 most common reef fish species for each MPA and comparison.  
 828

Species Name	NSC	ED	NF
<i>Balistes capriscus</i> (Gmelin, 1789)	x	x	x
<i>Calamus nodosus</i> (Randwall & Caldwell, 1966)	x	x	
<i>Centropristis striata</i> (Linnaeus, 1758)	x		x
<i>Cephalopholis cruentata</i> (Lacepède, 1802)		x	
<i>Cephalopholis fulva</i> (Linnaeus, 1758)	x		
<i>Epinephelus adscensionis</i> (Osbeck, 1765)	x		
<i>Epinephelus drummondhayi</i> Goode & Bean, 1878			x
<i>Epinephelus morio</i> (Valenciennes, 1828)	x	x	x
<i>Haemulon plumierii</i> (Lacepède, 1801)	x	x	
<i>Lutjanus campechanus</i> (Poey, 1860)		x	x
<i>Mycteroperca phenax</i> Jordan & Swain, 1884	x	x	x
<i>Pagrus pagrus</i> (Linnaeus, 1758)	x	x	x
<i>Rhomboplites aurorubens</i> (Cuvier, 1829)	x	x	x
<i>Seriola dumerili</i> (Risso, 1810)			x
<i>Seriola rivoliana</i> (Valenciennes, 1833)		x	x

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850 Table 3. ANOVA (type 3) results (F-statistics unless otherwise noted) on abundance ( $\log(x + 1)$ )  
851 transformed) for the top ten most abundant reef fish species (RFS) and for Red Porgy (RP).  
852 Global models include all MPA regions (“All”), as well as the results of sub-models for each  
853 MPA region: Northern South Carolina (NSC), Edisto (ED), North Florida (NF). F-statistics  
854 greater than 3.87 are significant ( $p < 0.05$ ) and in bold. Coefficient-Estimate ( $\beta$ ) is provided for  
855 the location by time interaction term.  
856

Model Type	Region	Location (Inside   Outside)	Time (Before   After)	Location by Time		Temperature	Depth	Relief	Substrate Density	Biota Density	Degrees of Freedom	Adjusted R <sup>2</sup>	F-statistic
				F	$\beta$								
<i>RFS</i>													
All	<b>4.90</b>	<b>12.5</b>	<b>4.0</b>	3.2	0.25	<b>24.3</b>	0.40	<b>10.4</b>	1.83	1.05	1185	0.06	<b>9.13</b>
NSC	-	<b>17.8</b>	3.21	0.76	0.16	<b>9.29</b>	<b>7.09</b>	2.00	0.13	0.21	505	0.12	<b>10.0</b>
ED	-	0.73	2.24	<b>4.86</b>	0.56	<b>16.6</b>	0.63	1.69	2.35	1.76	325	0.07	<b>43.92</b>
NF	-	<b>5.22</b>	-	-	-	<b>4.53</b>	0.21	2.55	0.26	0.31	341	0.05	<b>3.23</b>
<i>RP</i>													
All	<b>7.97</b>	1.5	<b>5.98</b>	0.23	-0.06	0.76	<b>4.10</b>	<b>18.1</b>	<b>4.03</b>	<b>6.3</b>	1185	0.08	<b>11.1</b>
NSC	-	0.19	3.47	0.55	-0.11	1.95	0.04	<b>12.4</b>	0.68	1.70	505	0.09	<b>7.07</b>
ED	-	<b>5.11</b>	0.69	0.66	0.19	0.05	1.88	0.42	3.40	<b>9.45</b>	325	0.12	<b>6.71</b>
NF	-	0.03	-	-	-	0.18	0.05	2.77	0.01	0.33	341	0.03	<b>2.55</b>

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877 **Figures**

878 Figure 1. Sampling Locations. Southeast Reef Fish Survey (SERFS) chevron trap sampling  
879 locations in the southeast United States (SEUS) Atlantic Ocean with the three Marine Protected  
880 Areas (MPA) and adjacent fished comparison areas analyzed in this study: Northern South  
881 Carolina, Edisto, and North Florida.  
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883 Figure 2. MPA Global Analysis of Changes in Reef Fish Abundances. Mean ( $\log(x + 1)$ ) +/-  
884 standard error (se) and linear model predicted values +/- se (red) for abundance of the ten most  
885 commonly caught managed reef fish species (upper panel) and Red Porgy (lower panel)  
886 inside/outside of three SEUS MPAs and before/after MPA implementation. Refer to Table 3 for  
887 summary of ANOVA (type 3) results. Values are based on trap data collected by the Southeast  
888 Reef Fish Survey 2000-2018, excluding samples collected in 2009 when MPAs were established.  
889

890 Figure 3. MPA Analysis of Changes in Reef Fish Abundances. Mean ( $\log(x + 1)$ ) +/- standard  
891 error of the ten most commonly caught managed reef fish species inside and outside of three  
892 SEUS MPAs Northern South Carolina (a), Edisto (b) and North Florida (c). Values are based on  
893 trap data collected by the Southeast Reef Fish Survey 2000-2018, excluding samples collected in  
894 2009 (vertical dashed line) when MPAs were established. Sample sizes (N) for each three-year  
895 period are reported below each graph (inside|outside).  
896

897 Figure 4. MPA Analysis of Changes in Red Porgy Abundances. Mean ( $\log(x+1)$ ) +/- standard  
898 error of the ten most commonly caught managed reef fish species inside and outside of three  
899 SEUS MPAs Northern South Carolina (a), Edisto (b) and North Florida (c). Values are based on  
900 trap data collected by the Southeast Reef Fish Survey 2000-2018, excluding samples collected in  
901 2009 (vertical dashed line) when MPAs were established. Sample sizes (N) for each three-year  
902 period are reported below each graph (inside|outside).  
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904 Figure 5. MPA Analysis of Changes in Reef Fish Sizes. Total length distributions (cm) of the ten  
905 most commonly caught managed reef fish species inside and outside three MPAs: Northern  
906 South Carolina (a), Edisto (b), North Florida (c). Values are based on trap data collected by the  
907 Southeast Reef Fish Survey 2000-2018, excluding samples collected in 2009 (vertical dashed  
908 line) when MPAs were established. One-sided t-test (assuming unequal variance) of inside-  
909 outside comparison used to assess significance. ( $p < 0.05 = *$ ,  $p < 0.01 = **$ ,  $p < 0.001 = ***$ ).  
910 Sample sizes (N) for each four-year period are reported below each graph (inside|outside).  
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912 Figure 6. MPA Analysis of Changes in Red Porgy Sizes. Total length distributions (cm) of Red  
913 Porgy inside and outside three MPAs: Northern South Carolina (a), Edisto (b), North Florida (c).  
914 Values are based on trap data collected by the Southeast Reef Fish Survey 2000-2018, excluding  
915 samples collected in 2009 (vertical dashed line) when MPAs were established. One-sided  
916 Wilcoxon rank sum test of inside-outside comparison used to assess significance. ( $p < 0.05 = *$ ,



917 p<0.01 = \*\*, p<0.001 = \*\*\*). Sample sizes (N) for each four-year period are reported below  
918 each graph (inside|outside).

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920 Figure 7. MPA Analysis of Changes in Percent of Fully-Recruited Red Porgy. Percent of fully-  
921 recruited Red Porgy (age 4 and above), three MPAs: Northern South Carolina (a), Edisto (b),  
922 North Florida (c). Values are based on trap data collected by the Southeast Reef Fish Survey  
923 2000-2018, excluding samples collected in 2009 (vertical dashed line) when MPAs were  
924 established. Significance tested for inside-outside comparisons using chi-square test (p<0.05 = \*,  
925 p<0.01 = \*\*, p<0.001 = \*\*\*). Sample sizes (N) for each four-year period are reported below  
926 each graph (inside|outside).

927