

1 **Hierarchical Modeling Assessment of the Influence of Watershed**
2 **Stressors on Fish and Invertebrate Species in Gulf of Mexico Estuaries**

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29

30 **Abstract**

31 The northern Gulf of Mexico (GoM) spans five U.S. states and encompasses estuaries that vary
32 greatly in size, shape, upstream river input, eutrophication status, and biotic communities. Given
33 the variability among these estuaries, assessing their biological condition relative to
34 anthropogenic stressors is challenging, but important to regional fisheries management and
35 habitat conservation initiatives. Here, a hierarchical generalized linear modeling approach was
36 developed to predict species presence in bottom trawl samples, using data from 33 estuaries over
37 a nineteen-year study period. This is the first GoM estuary assessment to leverage Gulf-wide
38 trawl data to develop species-level indicators and a quantitative index of estuary disturbance.
39 After controlling for sources of variability at the sampling event, estuary, state, and sampling
40 program levels, our approach screened for statistically significant relationships between
41 watershed-level anthropogenic stressors and fish and invertebrate species presence. Modeling
42 results indicate species level indicators with sensitivities to landscape stressor gradients. The
43 most influential stressors include total anthropogenic land use, crop land use, and the number of
44 toxic release sites in upstream watersheds, as well as agriculture in the shoreline buffer, each of
45 which was significantly related to between 21% and 39% of the 57 species studied. Averaging
46 the effects of these influential stressors across species, we develop a quantitative estuary stress
47 index that can be compared against benchmark conditions. In general, disturbance levels were
48 greatest in estuaries west of the Mississippi delta and in highly developed estuaries in southwest
49 Florida. Estuaries from the Florida panhandle to the eastern Mississippi delta had less
50 anthropogenic stress.

51 **Keywords:** biological assessment, species indicators, anthropogenic stressors, hierarchical
52 modeling, watershed development, Gulf of Mexico

53 1. Introduction

54 Fishing is central to the social and economic well-being of the northern Gulf of Mexico (GoM)
55 region of the United States (U.S.), making sustainable management of fisheries a regional
56 priority. The seafood industry in the Gulf States of Florida, Alabama, Mississippi, Louisiana, and
57 Texas (FL, AL, MS, LA, TX) contributed \$7.9 B to the 2012 U.S. Gross Domestic Product
58 (GDP) and provided 160,000 jobs to coastal residents (NMFS 2014), while recreational fisheries
59 provided an additional \$7.8 B to the regional GDP in 2012 as a result of the activities of 3.1 M
60 anglers (NMFS 2014). Some of the most valuable species in both the commercial and
61 recreational fisheries, including shrimps (Family: Penaeidae), Gulf Menhaden (*Brevoortia*
62 *patronus*), and Spotted Seatrout (*Cynoscion nebulosus*), have strong affiliations to estuary
63 habitats for a portion of their life cycles. The estuaries of the GoM are subject to disturbance
64 from a wide range of anthropogenic activities, potentially putting commercial and recreational
65 fisheries at risk. Understanding the spatial patterns and causes of degradation to estuary fisheries
66 and fish habitats are important science priorities.

67 Environmental degradation in some areas of the GoM is already advanced due to anthropogenic
68 disturbances that include hydrologic alteration, eutrophication, toxic pollution, and overfishing
69 (NRC, 2000; Rabalais et al., 2002; Yáñez-Arancibia and Day, 2004, Howarth and Marino, 2006).
70 Altered patterns of freshwater inflow to GoM estuaries have resulted from upstream damming,
71 river channelization, and water abstraction (Harwell, 1997; Flannery et al., 2002), which have led
72 to changed salinity regimes, reduced dilution of estuary pollutants, and land subsidence (Day et
73 al., 2000). Excessive runoff of nitrogen and phosphorus from fertilizers and urban activities in
74 catchments has caused eutrophication, and in severe cases, low dissolved oxygen and fish kills
75 (Rabalais et al., 2002; Diaz and Rosenberg, 2008; 2011). Toxic chemicals associated with
76 industry, urban development, and agriculture are strongly concentrated in some areas and have
77 been shown to negatively affect benthic organisms in the GoM (Brown et al., 2000). Some of
78 these toxic releases originate from the petroleum industry, which is especially concentrated in
79 coastal areas of LA and eastern TX (Adams et al., 2004). Estuarine fish and invertebrate species
80 integrate habitat conditions differently over both time and space and can be helpful as biological
81 indicators of larger trends in ecosystem degradation (e.g., Macauley et al. 1999).

82 A central challenge in biological assessment is to distinguish the responses of species to
83 anthropogenic stress from the high levels of natural background variation common to ecological
84 systems (Hawkins et al., 2010). This challenge is particularly acute in GoM estuaries where the
85 daily and seasonal variability in temperature, salinity, and dissolved oxygen leads to annual
86 variation in biological community composition and structure (Peterson and Ross, 1991; Akin et
87 al., 2003; Baltz et al., 1993; Gelwick et al., 2001; Granados-Dieseldorff and Baltz, 2008). While
88 high natural environmental variability can be stressful to many species, anthropogenic stress to
89 estuaries has been shown to result in dominance by opportunistic habitat or trophic generalists, to
90 the detriment of rare or specialized taxa (Felley, 1987; Chesney and Baltz, 2001; Lewis et al.,
91 2011). Such findings suggest that some species may be particularly sensitive to human impacts,
92 including even estuarine species adapted to high degrees of natural environmental variation.
93 Many estuarine studies conducted to date have focused on responses by groups of species with
94 similar life history or functional characteristics (i.e. community metrics; Macauley et al., 1999;
95 Summers, 2001; Hughes et al., 2002; Meng et al., 2002; Jordan et al., 2010; Cabral et al., 2012).

96 However, evidence from freshwater systems suggests that community metrics may obscure
97 biological responses to stressor gradients that can be detected in species-specific indicators
98 (Baker and King, 2010; King and Baker, 2010). Further, individual species indicators may also
99 allow for more direct linkages to management by focusing on populations of economically
100 valuable taxa or taxa with high conservation value.

101 Multiple approaches have been used to account for natural background variation in biological
102 assessment. One approach is to define different biological indicators within discrete salinity
103 zones (Coates et al., 2007; Briene et al., 2010; Cabral et al., 2012) or different types of estuaries
104 (Harrison and Whitfield, 2006). Another approach is to screen for indicators that are sensitive to
105 anthropogenic stress, but insensitive to natural gradients (Jordan et al., 2010). Moreover, models
106 can be developed and used to account for the effects of natural variables before testing for
107 indicator sensitivity (Engle et al., 1994). Multivariable models (e.g. multiple linear regression)
108 have proven useful for predicting species or community responses to both natural and
109 anthropogenic gradients (Lewis et al., 2007; Courrat et al., 2009; Delpech et al., 2010). By
110 controlling for natural variation with model coefficients, biological responses to one or multiple
111 ecological stressors can be predicted at different stressor levels which can be particularly helpful
112 for judging whether current conditions differ from a benchmark or reference condition (Hawkins
113 et al., 2010). Without benchmarks, little context exists for interpreting the measured value of an
114 ecological resource, which can vary substantially with natural differences among sites.

115
116 Efforts to classify the ecological status of estuaries have occurred globally over the past two
117 decades. In the U.S., studies have focused on characterizing estuaries based on their water
118 quality, susceptibility to pollution based on geomorphologic and flow conditions, and watershed
119 stressors such as land cover and point sources of pollution (Bricker et al., 2008; Greene et al.,
120 2015). Some studies have classified U.S. estuaries based on their fish populations (Gleason et al.,
121 2011; Hughes et al., 2014), but only a few nekton species in limited regions have been modeled
122 to connect fish presence to estuary anthropogenic stress (Toft et al., 2015). In Europe, regional
123 and country specific multi-metric indices have been developed based on biological communities
124 to calculate overall ecological health (Breine et al., 2007; Coates et al., 2007; Delpech et al.,
125 2010; Cabral et al., 2012; Harrison and Kelly, 2013), but limited progress has been made toward
126 showing strong relationships between these indices and anthropogenic stressors (Pasquad et al.,
127 2013). Such analyses have been complicated by the additional effort required to “intercalibrate”
128 the results from different studies in order to make intra-continental comparisons. In particular,
129 setting regional benchmark conditions and comparing studies with different sampling protocols
130 or metrics is an ongoing challenge not easily resolved (Poikane et al., 2014). Though much
131 progress has been made in sampling and quantifying estuary ecological status, further efforts and
132 techniques are needed to model species and biological communities across varying natural
133 settings, link biological conditions to watershed stressors, and set benchmark conditions.

134
135 Hierarchical (or ‘multi-level’) modeling provides a potential methodology for linking watershed
136 stressors and estuary biological condition, accounting for discrepancies among sampling
137 programs and natural variability among estuaries. Hierarchical modeling accounts for variability
138 among different groups of data at different spatial or organizational levels using regression
139 coefficients (i.e., ‘random effects’) that vary by group as members of a common statistical
140 hyperdistribution (Gelman and Hill, 2006). This extension of classical regression modeling

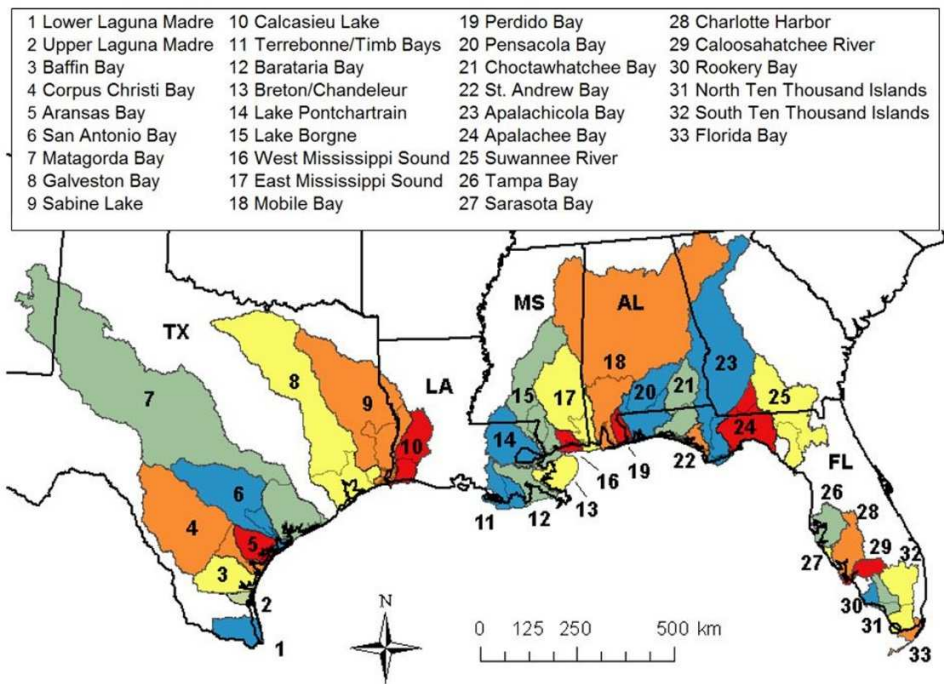
141 accounts for intra-class correlation among data from common groups (i.e., estuaries, states, trawl
142 programs), allowing for statistically valid hypothesis testing of group-level predictor variables
143 (Gelman et al., 2014). Thus, for grouped data, hierarchical modeling is often an improvement
144 over classical regression modeling in terms of both predictive performance and causal inference
145 (Wikle, 2003a; Cressie et al., 2009; Qian et al., 2010), and these models have been used
146 extensively to study environmental and ecological systems (Wikle et al., 1998; Wikle, 2003b;
147 Clark and Gelfand, 2006; Bolker et al., 2009; Kashuba et al., 2010; Cuffney et al., 2011). In this
148 study, hierarchical modeling is instrumental in controlling for the variability of species presence
149 across different estuaries, states, and monitoring programs.

150 The aim of the current study is to identify key sources of watershed stress (i.e., stressors) that are
151 related to species presence in GoM estuaries, and to aggregate these relationships to assess the
152 relative intensity of watershed stress in each estuary using hierarchical generalized linear
153 modeling. This new approach to assessing the biological health of estuaries allows us to: (1)
154 combine nearly 70,000 trawl samples collected by separate research efforts; (2) control for
155 natural environmental variation while identifying statistically significant estuary-level stressors
156 affecting the presence of fish and invertebrate species; and (3) create an estuary stress index that
157 quantifies the amount of anthropogenic stress affecting GoM estuaries as compared to
158 benchmark conditions in the region.

159 **2. Methods**

160 *2.1 Study Area*

161 Our study spanned 33 estuaries across the five U.S. GoM states (Fig. 1). Estuary habitats were
162 classified to include open water and wetland classes from the Coastal Change Analysis Program
163 (C-CAP) dataset (NOAA, 2006) and the National Wetlands Inventory (NWI; USFWS, 2012). To
164 summarize landscape influence on each estuary, it was necessary to define their spatial extents
165 and tributary influences. The seaward extent of estuaries was limited to the 4-m depth contour
166 based on an examination of plots of salinity-at-depth. The salinities at the defined seaward depth
167 did not generally drop below 32 practical salinity units (psu). The landward extent of the
168 estuaries was drawn to include open water areas as well as estuarine emergent wetlands from C-
169 CAP. Atchafalaya Bay, LA was excluded because it is a statistical outlier in terms of its
170 freshwater inflow and nitrogen load (from the greater Mississippi River Basin), and thus difficult
171 to compare to other GoM estuaries. Additional coastal areas were omitted from this analysis
172 because they did not meet the conventional definition of an estuary (i.e., a partially enclosed
173 coastal body of brackish water with one or more rivers or streams flowing into it and with a free
174 connection to the open sea; Pritchard, 1967) or they lacked sufficient watershed level stressor
175 data needed for analysis. One estuary, Rookery Bay, was included in the final analysis because it
176 had watershed level stressor data even though it did not possess any biological data (i.e., trawl
177 data).



178

179 **Fig. 1** Basins for the 33 estuaries and estuarine drainage areas (EDAs), representing the most
 180 downstream portion (HUC 8) of each basin, are delineated using gray lines while state lines are
 181 shown in black. For some smaller drainage basins, the total basin and EDA are equivalent (no
 182 gray line within the colored basin). The GoM includes five states: Texas (TX), Louisiana (LA),
 183 Mississippi (MS), Alabama (AL), and Florida (FL).

184

185 Data were collected at two different spatio-temporal scales. Throughout this paper, “event-level
 186 data” refer to the data recorded at the specific times and locations of individual bottom trawl
 187 samples. “Estuary-level data” refer to average conditions within estuaries or in the watersheds
 188 connecting to an estuary. Physical features include both event-level data (temperature, salinity
 189 and distance-to-shore) and estuary-level data such as estuary volume, estuary area, percent of
 190 estuary open to the sea, and average freshwater inflows. All anthropogenic stressors included in
 191 this analysis, such as toxic releases and land covers, are estuary-level data.

192 *2.2 Trawl Events*

193 Sampling events, performed by state and federal programs, consisted of towing otter trawls
 194 through estuary bottom waters for set distances or periods of time to determine the presence and
 195 abundance of different fish and invertebrates living in those areas. This study analyzed 69,570
 196 bottom trawl samples from 1991 through 2009 (Table 1) gathered by the five GoM states (FL,
 197 AL, MS, LA, TX) and two federal research programs run by the U.S. Environmental Protection
 198 Agency (EPA): the Environmental Monitoring and Assessment Program (EMAP) and National
 199 Coastal Assessment (NCA). Sampling gears and protocols varied across programs (Table 1). For

200 instance, programs used nets with varying dimensions and cod-end mesh sizes, and towed for
 201 different distances and at different speeds. Therefore, a measure of effort (hectares (ha) sampled)
 202 was derived for each sampling event, and the cod-end mesh size was recorded for each sampling
 203 program. Variation that existed in sampling protocols was accounted for by a hierarchical
 204 grouping of trawl data that allowed the model intercept to vary by sampling program (see section
 205 2.6).

206 **Table 1** Summary of fish and invertebrate trawl data for
 207 Florida (FL), Alabama (AL), Mississippi (MS),
 208 Louisiana (LA), Texas (TX), Environmental Monitoring
 209 and Assessment Program (EMAP), and National Coastal
 210 Assessment (NCA).

Program	# of trawls	# of species recorded	Mesh size (mm)	Trawl effort (ha)	Time period
FL	9,580	213	3.2	5.6	1991-2005
AL	2,620	102	9.5	4.2	1991-2006
MS	708	25	6.4	4.2	1991-2005
LA	23,580	209	6.4	3.2	1991-2007
TX	31,870	488	38	2.8	1991-2009
EMAP	418	157	25	5.0	1991-1994
NCA	795	244	38	5.0	2000-2004

211

212 For each bottom trawl event, fish and invertebrate species were identified and enumerated along
 213 with basic environmental data such as location, temperature, and salinity. Some trawl events
 214 yielded no data because no species were collected, while most events contained records for
 215 multiple species. Data on over 500,000 fish and invertebrate individuals collected from the trawl
 216 events were used in this study, but only species that were sampled by at least six of the seven
 217 monitoring programs and caught in a minimum of 120 trawls were used for final results (see
 218 section 2.7). This limited our study to 57 fish and invertebrate species (Table 2) whose ranges
 219 were widely distributed throughout the GoM and had sufficient data to adequately parameterize
 220 logistic models (Peduzzi et al., 1996; Allison, 2012).

221 *2.3 Event-level Variables*

222 Predictor variables were compiled at both the event and estuary levels. Preliminary analyses
 223 suggested that temperature, salinity and distance-to-shore were important event-level variables
 224 for the majority of species while average estuary depth was not. Temperature and salinity can
 225 determine important characteristics of fish habitat due to their effect on primary production and
 226 fish metabolism (Fry, 1971; Brett and Groves, 1979), while distance-to-shore can act as a proxy
 227 for near-shore habitats and changes in water depth. Reliable event-level depth estimates were
 228 missing for a substantial portion of the trawl data and as such this variable had to be excluded
 229 from the analysis, despite its obvious importance for fish habitat selection and trawl catch
 230 efficiency. Temperature, salinity, and distance-to-shore exhibited substantial variability as seen
 231 in Table 3. The variations in the event-level predictors reflect, to a large extent, the natural

232 heterogeneity of estuarine conditions across the study area. While hydrologic alteration in the
 233 GoM can affect freshwater inflows and estuary salinity levels (Orlando et al., 1993; Day et al.,
 234 2000), these large-scale anthropogenic modifications were not considered directly in this study
 235 because they could not be explicitly quantified, though they are likely related to some of the
 236 available watershed stressor variables.

237

238 **Table 2** Modeled fish and invertebrate species along with overall percent presence in trawl
 239 samples (T) and in estuaries (E).

Common/Scientific species name	% pres.		Common/Scientific species name	% pres.	
	T	E		T	E
Atl. Croaker/ <i>Micropogonias undulatus</i>	52	85	Southern Flounder/ <i>Paralichthys lethostigma</i>	5	73
Bay Anchovy/ <i>Anchoa mitchilli</i>	49	85	Striped Mullet/ <i>Mugil cephalus</i>	5	61
Blue Crab / <i>Callinectes sapidus</i>	42	94	Black Drum/ <i>Pogonias cromis</i>	5	64
Spot/ <i>Leiostomus xanthurus</i>	39	82	Striped Anchovy/ <i>Anchoa hepsetus</i>	4	76
White Shrimp/ <i>Litopenaeus setiferus</i>	35	70	Silver Jenny/ <i>Eucinostomus gula</i>	3	82
Brown Shrimp/ <i>Farfantepenaeus aztecus</i>	34	70	Lookdown/ <i>Selene vomer</i>	3	73
Hardhead Catfish/ <i>Ariopsis felis</i>	30	85	Star Drum/ <i>Stellifer lanceolatus</i>	3	52
Sand Seatrout/ <i>Cynoscion arenarius</i>	28	85	Gulf Toadfish/ <i>Opsanus beta</i>	3	76
Pinfish/ <i>Lagodon rhomboides</i>	27	94	Lined Sole/ <i>Achirus lineatus</i>	3	73
Gulf Menhaden/ <i>Brevoortia patronus</i>	20	79	Atl. Moonfish/ <i>Selene setapinnis</i>	3	55
Silver Perch/ <i>Bairdiella chrysoura</i>	20	82	Chain Pipefish/ <i>Syngnathus louisianae</i>	2	76
Pink Shrimp/ <i>Farfantepenaeus duorarum</i>	13	85	Crevalle Jack/ <i>Caranx hippos</i>	2	52
Least Puffer/ <i>Sphoeroides parvus</i>	12	70	Silver Seatrout/ <i>Cynoscion nothus</i>	2	70
Bay Whiff/ <i>Citharichthys spilopterus</i>	11	67	Scaled Sardine/ <i>Harengula jaguana</i>	2	55
Gafftopsail Catfish/ <i>Bagre marinus</i>	10	79	Blue Catfish/ <i>Ictalurus furcatus</i>	2	67
Black Tonguefish/ <i>Symphurus plagiusa</i>	9	76	Gizzard Shad/ <i>Dorosoma cepedianum</i>	2	64
Fringed Flounder/ <i>Etropus crossotus</i>	9	79	Spotfin Mojarra/ <i>Eucinostomus argenteus</i>	2	52
Atl. stingray/ <i>Dasyatis sabina</i>	8	73	Gulf Pipefish/ <i>Syngnathus scovelli</i>	2	79
Inshore Lizardfish/ <i>Synodus foetens</i>	8	97	Naked Goby/ <i>Gobiosoma bosc</i>	1	73
Hog Choker/ <i>Trinectes maculatus</i>	7	85	Lane Snapper/ <i>Lutjanus synagris</i>	1	58
Bighead Searobin/ <i>Prionotus tribulus</i>	7	85	Red Drum/ <i>Sciaenops ocellatus</i>	1	64
Atl. Bumper/ <i>Chloroscombrus chrysurus</i>	7	76	Atl. Threadfin Herring/ <i>Opisthonema oglinum</i>	1	55
Atl. Spadefish/ <i>Chetodipterus faber</i>	7	73	White Mullet/ <i>Mugil curema</i>	1	48
Ground Mullet/ <i>Menticirrhus americanus</i>	7	82	Bluntnose Jack/ <i>Hemicaranx amblyrhynchus</i>	0.9	58
Pigfish/ <i>Orthopristis chrysoptera</i>	7	82	Spanish Mackerel/ <i>Scomberomorus maculatus</i>	0.8	61
Gulf Butterfish/ <i>Peprilus burti</i>	6	70	Ladyfish/ <i>Elops saurus</i>	0.8	64
Spotted Seatrout/ <i>Cynoscion nebulosus</i>	6	79	Lined Seahorse/ <i>Hippocampus erectus</i>	0.6	48
Threadfin Shad/ <i>Dorosoma petenense</i>	6	61	Sea Bass/ <i>Centropristis philadelphica</i>	0.3	52
Atl. Cutlassfish/ <i>Trichiurus lepturus</i>	5	85			

240

241 **Table 3.** Summary of event-level data for trawl events.

	Temperature (°C)	Salinity (psu)	Distance-to- shore (km)
Mean	23.1	18.0	2.7
Standard deviation	6.1	11.0	3.3
2.5% quantile	11	0	0.04
97.5% quantile	31	38	14.1

242

243 All data were checked for potential outliers. Events that reported temperature values over 35° C
 244 or below 6° C were considered suspect and removed. Similarly, all non-winter records with
 245 temperatures below 10° C were excluded. In total, 285 trawl events were removed for having
 246 temperature values outside of these ranges, representing 0.4% of the original dataset. Given that
 247 salinity values above 40 psu are common in hyper-saline estuaries such as Baffin Bay, TX, no
 248 salinity values were excluded from this analysis (maximum salinity = 66 psu). Sample months
 249 were aggregated into seasons: spring (March, April, May), summer (June, July, August), fall
 250 (September, October, November), and winter (December, January, February), which were
 251 considered as a categorical variable within the models.

252 *2.4 Estuary-level Variables*

253 Land cover data were compiled and analyzed at three different spatial scales: basin, estuarine
 254 drainage area (EDA; Fig. 1) and shoreline buffer. Basin data covers were summarized from the
 255 entire drainage area of the estuary to the topographic divide with adjacent river basins with
 256 outlets to the sea, while EDAs were limited only to the lowest most proximate 8-digit hydrologic
 257 unit code (HUC) watershed that drains to the coast (NOAA/NOS, 1985). The shoreline buffer
 258 was defined as the 500-meter buffer inland from each estuary boundary. EDA sizes ranged from
 259 246 km² to 13,690 km², while basin areas ranged from 330 km² to 121,762 km². For some
 260 estuaries with entirely coastal drainage areas, basin and EDA values were the same (e.g.,
 261 Sarasota Bay, FL, Perdido Bay, AL, and Baffin Bay, TX; Fig 1). Land cover data from 1992,
 262 2001 and 2006 were highly correlated ($r > .98$) such that all estuary-level land cover data was
 263 calculated using the 2001 National Land Cover Database (Homer et al., 2007) representing the
 264 middle of the study period and supplemented with wetland classifications from C-CAP (NOAA,
 265 2006) and the National Wetlands Inventory (USFWS, 2012). We grouped land cover categories
 266 to reduce potential collinearity and to generate combinations of land cover types expected to
 267 have similar impacts on estuary habitat quality. The “Agriculture” class was formed by grouping
 268 crop and pasture, both of which are expected to yield excess nutrients to waterways while the
 269 “Anthropogenic” class aggregated urban, crop, and pasture into one class to represent land use
 270 predominantly affected by humans. The land cover classes adopted in the model were
 271 normalized by the total land area (A_L) of the given spatial unit (basin, EDA, or shoreline buffer),
 272 creating percent land cover values for each spatial scale. Other approaches to normalizing land
 273 cover are described below.

274 Watershed land cover, population, number of toxic release sites, and number of National
 275 Pollutant Discharge Elimination System (NPDES) sites (USEPA, 2015) can be interpreted as
 276 proxies for pollution, both at the basin and EDA levels. For example, crop areas may export
 277 nitrogen, phosphorous, pesticides, and various other agricultural chemicals downstream to
 278 estuaries. We would have preferred to use more direct predictors like nitrogen, phosphorus, or
 279 other pollutant loads, but reliable loading data were unavailable for many estuaries. Land-based
 280 pollution loads are often attenuated through biogeochemical processing, settling, and flushing. In
 281 the current study, it was infeasible to estimate reaction rates or settling velocities for the diverse
 282 range of potentially important pollutants. In an effort to indirectly account for settling and
 283 flushing, many estuary-level variables were normalized by estuary area (A_E) and flow (Q), in
 284 addition to A_L , creating several stressors per variable (e.g., EDA Crop/ A_E , EDA Crop/ Q , and
 285 EDA Crop/ A_L ; Table 4). Shoreline buffers are not large enough to produce significant loads, but
 286 may affect nearshore habitat conditions, and were thus normalized by only A_E . Five estuarine
 287 variables from the 2007 National Estuarine Eutrophication Assessment (Bricker et al., 2008),
 288 including estuary average salinity, estuary percent open to sea (%), hypoxic condition, and toxic
 289 algal condition (scaled from 0 = no problem to 3 = high), and eutrophication condition (scaled
 290 from 1 = low to 5 = high), were also included without normalization. In total, 47 candidate
 291 estuary-level variables were considered (Table 4). Basic estuary-level data can be found in
 292 Supplementary Material, SM 1 and SM 2.

293 **Table 4** Estuary-level predictor variables considered in the modeling analysis using various
 294 normalization options (1 = no normalization; A_E = estuary area (km²); Q = flow (m³/day); A_L =
 295 total land area). Note: I(0:3) indicates an integer variable (0, 1, 2, or 3).

Variable	Unit	1	A_E	Q	A_L
Watershed					
Shoreline Urban	km ²		X		X
Shoreline Crop	km ²		X		X
Shoreline Agriculture	km ²		X		X
Shoreline Anthropogenic	km ²		X		X
Shoreline Wetlands	km ²		X		X
EDA Urban	km ²		X	X	X
EDA Crop	km ²		X	X	X
EDA Agriculture	km ²		X	X	X
EDA Anthropogenic	km ²		X	X	X
Basin Urban	km ²		X	X	X
Basin Crop	km ²		X	X	X
Basin Agriculture	km ²		X	X	X
Basin Anthropogenic	km ²		X	X	X
EDA toxic releases	#		X	X	
EDA NPDES	#		X	X	
EDA population	#		X	X	
Basin population	#		X	X	
Estuary					
Mean estuary salinity	psu	X			
Estuary percent open to sea	%	X			
Hypoxic condition	I(0:3)	X			
Toxic algal condition	I(0:3)	X			
Eutrophic condition	I(1:5)	X			

296 2.5 Hierarchical Generalized Linear Model

297 Species presence and absence in trawl samples was modeled as a binary response, where trawls
298 that contained a given species were assigned a value of one and trawls that did not collect that
299 species were assigned a value of zero. A common approach for modeling binary data is to use a
300 generalized linear model that extends the framework of linear regression modeling to non-
301 normally distributed variables by the use of a link function. These types of models are used
302 extensively in many fields, including ecology (Guisan and Zimmermann, 2000; Guisan et al.,
303 2002; Gelfand et al., 2005; Gelman and Hill, 2006; Latimer et al., 2006, Bolker, 2008).

$$304 \quad P(y_i) = \text{logit}^{-1}(\mathbf{X}_i \boldsymbol{\beta}) \quad (\text{Eq. 1})$$

305 where $P(y_i)$ is the probability of fish presence in trawl i , \mathbf{X}_i is a matrix of predictor variables, and
306 $\boldsymbol{\beta}$ is a vector of corresponding model coefficients. The inverse-logit function is used to back
307 transform the response from the continuous modeling domain ($-\infty$ to $+\infty$) to probability space
308 values between 0 and 1: $\text{logit}^{-1}(x) = \frac{e^x}{1+e^x}$.

309 In addition to accounting for binary responses, our model is constructed hierarchically in order to
310 capture different levels of organization within the data. Hierarchical models use “random
311 effects” to account for data that are similar and can be grouped as members of a common
312 statistical hyperdistribution (Gelman and Hill, 2006). This approach helps account for intra-class
313 correlation among grouped data allowing for statistically valid hypothesis testing of group-level
314 (i.e., estuary-level) predictor variables (Gelman et al., 2014). Random effects were used in this
315 study to account for variation at the estuary, program, and state level that was not explained by
316 other predictors in the model. Without this added model flexibility, prescreening would not have
317 been possible without much noise from spurious correlations between predictors and watershed
318 stressors (see Section 4.4).

319 2.6 Single-stressor Models

320 For each fish and invertebrate species, we initially screened all 47 possible estuary-level
321 predictor variables (x_{pred}) separately while controlling for event-level variables. Logistic
322 hierarchical models were fit using the “lme4” R package with parallel processing provided by the
323 “doparallel” R package (R Development Core Team, 2011; Bates et al., 2013; Calaway et al.,
324 2015). The adopted model structure is shown below.

$$325 \quad P(y) = \text{logit}^{-1}(\beta_0 + \beta_{\text{season}} + \alpha_{\text{estuary}} + \alpha_{\text{state}} + \alpha_{\text{program}} + \beta_{\text{temp}} * x_{\text{temp}} + \beta_{\text{sal}} * x_{\text{sal}} + \beta_{\text{salsq}} * x_{\text{salsq}} + \\ 326 \quad \beta_{\text{dist}} * x_{\text{dist}} + \beta_{\text{pred}} * x_{\text{pred}}) \quad (\text{Eq. 2})$$

327 where $P(y)$ is the probability of occurrence of a given species, β_0 is the overall y-intercept for the
328 model, and β_{season} is a categorical variable used to account for seasonal differences. Random
329 effects, α_{estuary} , α_{state} , and α_{program} , were assumed to follow normally distributed hyperdistributions
330 centered around zero. For each species modeled, 32 different values of α_{estuary} were returned
331 (Rookery Bay is excluded due to no trawl events), five values for α_{state} and seven values for
332 α_{program} . Slope coefficients β_{temp} , β_{sal} , β_{salsq} , and β_{dist} were determined for the event-level effects of
333 temperature (x_{temp}), salinity (x_{sal}), salinity² (x_{salsq}), and distance-to-shore (x_{dist}), respectively. The

334 square of salinity (x_{salsq}) was included because preliminary analysis indicated a potential non-
335 linear (quadratic) relationship between salinity and probability of fish or invertebrate presence.
336 β_{pred} is the slope coefficient of the estuary-level anthropogenic variable being screened. These
337 single-stressor models were used to test the significance of each estuary-level predictor (Table 4)
338 on the occurrence of every fish and invertebrate species (Table 2). Predictor variables were
339 considered statistically significant when their corresponding β_{pred} coefficients were significantly
340 different from zero at a 95% confidence level ($p < 0.05$).

341 *2.7 Multi-stressor Models*

342 Candidate estuary-level variables for multi-stressor modeling were selected based on the single
343 stressor modeling results. Favoring parsimonious multi-stressor models, we selected only
344 estuary-level variables that were found to be significant for over 20% of the species.
345 Furthermore, variables were tested for multicollinearity, and when one or more variables were
346 found to be correlated ($|r| \geq 0.65$; Supplementary Material, SM 3), only the most significant
347 variable was included. This process reflects two assumptions: 1) stressor variables with truly
348 mechanistic underpinnings should affect a broad range of species across the GoM and 2)
349 variables that represented similar mechanisms (e.g., Basin Crop/ A_L is a subset of Basin
350 Agriculture/ A_L) should only enter the model once. Performing variable selection on a subset of
351 probable stressors instead of the entire 47 dependent variables (Table 4) avoids the inclusion of
352 spurious or “noisy” predictors and thus leads to more robust models (Cohen, 1990; Derksen and
353 Kesselman, 1992).

354 For each species, all event-level variables (temperature, salinity, salinity², and distance to shore)
355 and the selected candidate estuary-level variables were included together in a backward model
356 selection procedure using the “LMERConvenienceFunctions” R package (Tremblay and Ransijn,
357 2015). Backward selection eliminates variables in a step-wise fashion based on statistical
358 significance, until the final model contains only predictors significant at the 95% confidence
359 level. Note that since multi-stressor models could have up to nine predictors (four natural
360 variables and five anthropogenic ones; Eq. 2 and Table 5) and three levels of random effects
361 based on preliminary modeling results, these models were only developed for species caught in
362 at least 120 trawl events. This insures that we have at least 10 observations for each possible
363 predictor and level of random effect (Peduzzi et al., 1996; Allison, 2012). Model selection using
364 Akaike Information Criterion (AIC) and Bayesian Information Criterion (BIC) was also explored
365 (Sakamoto et al., 1986; Schwarz, 1978), but both of these criteria added insignificant predictors
366 as well as eliminated significant predictors making these methods not desirable for the overall
367 project goals.

368

369

370

371 **Table 5** Summary of highly significant estuary-level stressor variables (statistically related to
 372 over 20% of species), as determined from the single-stressor models. Negative relationships
 373 indicate the percent of significant relationships that are negative. Candidate variables are those
 374 tested in the multi-stressor models.

Stressor	Species affected (%)	Negative relationships (%)	Candidate variables
Basin Anthropogenic/ A_L	53	77	X
Mean estuary salinity	49	61	X
Basin Agriculture/ A_L	47	70	
Basin Crop/ A_L	44	72	
Basin Crop/ Q	39	73	X
Basin Anthropogenic/ Q	37	67	
Shoreline Agriculture/ A_L	35	15	X
Basin Agriculture/ Q	35	70	
EDA toxic releases/ A_E	23	85	X
Basin Urban/ Q	21	58	

375
 376 *2.8 Assessment of Estuary Level Anthropogenic Stress*
 377 Once the multi-stressor models were developed, they were used to create an estuary stress index
 378 which quantifies the amount of anthropogenic disturbance in estuaries by comparing sampled
 379 conditions to those predicted under two different benchmarks of anthropogenic disturbance.
 380 Least disturbed conditions (LDC) were defined as the minimum value of each stressor observed
 381 at the regional scale (SM 1), and minimally disturbed conditions (MDC) were predicted by
 382 setting stressor values in the models to zero. LDC and MDC are benchmarks of ecological
 383 condition frequently used to assess the degree to which the current conditions of a particular
 384 system deviate from a more desirable state (Stoddard et al., 2006; Hawkins et al., 2010). We
 385 used the region-wide stressor minima to specify LDC as an example of one benchmark to assess
 386 estuaries in the GoM, but sub-regional LDCs might also be developed and used within this same
 387 framework. To judge the amount of anthropogenic watershed stress on each estuarine system, the
 388 absolute value of each stressor coefficient was averaged across all species to determine the mean
 389 absolute rate of change in probability of species presence (on the log-odds scale) associated with
 390 each stressor. Mean absolute coefficients were then multiplied by the actual estuary-level
 391 conditions and compared to both LDC and MDC to create the estuary stress index.

392 **3. Results**

393 *3.1 Single-stressor Models*

394 All estuary-level stressor variables (Table 4) were screened in single-stressor models (Eq. 2) that
 395 controlled for natural variability at the event-level (e.g., temperature, salinity) as well as for
 396 intra-class correlation and unknown variability at group levels (i.e., estuary, state, program).
 397 Variables were then ranked according to the number of species with which they had significant
 398 relationships (Table 5). Basin Anthropogenic/ A_L , Basin Crop/ A_L , and Basin Agriculture/ A_L were
 399 found to be statistically significant for over 40% of the assessed fish and invertebrate species
 400 with more than 70% of those significant relationships being negative. Of the other anthropogenic
 401 candidate stressors, Basin Crop/ Q and EDA toxic releases/ A_E were also negatively related to the

402 majority of the species, while Shoreline Agriculture/ A_L had a predominantly positive
403 relationship. Mean estuary salinity was found to be a significant predictor for 49% of modeled
404 species.

405 3.2 Multi-stressor Models

406 Multi-stressor models for individual fish and invertebrate species were developed by performing
407 backward variable selection on event-level variables (temp, sal, salsq, and dist) and the five
408 candidate estuary-level stressors identified in Table 5 (see Methods section for selection criteria).
409 This resulted in a unique multi-stressor model for each fish and invertebrate species (SM 4). For
410 example, the model for Silver Perch (*Bairdiella chrysoura*) was:

$$\begin{aligned} 411 \quad P(y) = \text{logit}^{-1} & (-2.92 + 0.02*[\text{temp}] + 0.04*[\text{sal}] - .001*[\text{salsq}] \\ 412 & - 0.05*[\text{dist}] - 0.40*[\text{EDA toxic releases}/A_E] - 0.22*[\text{Basin Crop}/Q] \\ 413 & + 0.29*[\text{Mean estuary salinity}] + \beta_{\text{season}} + \alpha_{\text{estuary}} + \alpha_{\text{state}} + \alpha_{\text{program}}) \quad (\text{Eq. 3}) \end{aligned}$$

414 The model indicates that Silver Perch presence increases with higher temperatures and in areas
415 closer to the shore. The relationship between salinity and Silver Perch is quadratic with an
416 optimal value at 20 psu, which is consistent with the fact that the Silver Perch prefer higher
417 saline estuaries. Silver Perch was negatively related to EDA toxic releases/ A_E and Basin Crop/ Q .
418 Note, coefficients for event-level variables are calculated in their natural units (i.e., °C) while
419 coefficients for estuary-level variables (e.g., EDA toxic releases/ A_E , estuary salinity) are based
420 on their normalized values (with mean of zero and standard deviation of one). Therefore, a
421 change of 1° C at the sampling spot has an effect of 0.02 on the log odds of Silver Perch presence
422 while the change of one standard deviation of EDA toxic releases/ A_E (6.3×10^{-4} releases/km²)
423 represents a change of 0.40 (SM 1,4). Multi-stressor models can be used to predict changes in
424 species presence within estuarine systems due to changes in estuary-level stressors (SM 5).

425 The fraction of anthropogenic (agricultural plus urban) land cover in an estuarine basin (Basin
426 Anthropogenic/ A_L) was found to be the most significant anthropogenic stressor in multi-stressor
427 models affecting 39% of species, of which 77% showed a negative relationship (Table 6). Basin
428 Crop/ Q was related to 37% of species, with 86% of those relationships being negative. Shoreline
429 Agriculture/ A_L was related to 23% of the species with most relationships being positive, and
430 EDA toxic releases/ A_E was related to 21% of the species with most relationships being negative.
431 Average estuary salinity was related to 25% of species as well. Mean absolute values of stressor
432 coefficients in log odds showed that Basin Anthropogenic/ A_L and Basin Crop/ Q exerted the
433 largest influence on species presence throughout the GoM (Table 6).

434

435

436

437 **Table 6** Percent of species significantly related to candidate estuary-level stressors in multi-
438 stressor models and percent of those relationships that are negative. Mean absolute change is
439 calculated by averaging the absolute value of the slope coefficients for all 57 species.

Stressor	Species affected (%)	Negative relationships (%)	Mean absolute change (log odds)
Basin Anthropogenic/ A_L	39	77	.27
Basin Crop/ Q	37	86	.24
Shoreline Agriculture/ A_L	23	8	.15
EDA toxic releases/ A_E	21	83	.14
Mean estuary salinity	25	36	.16

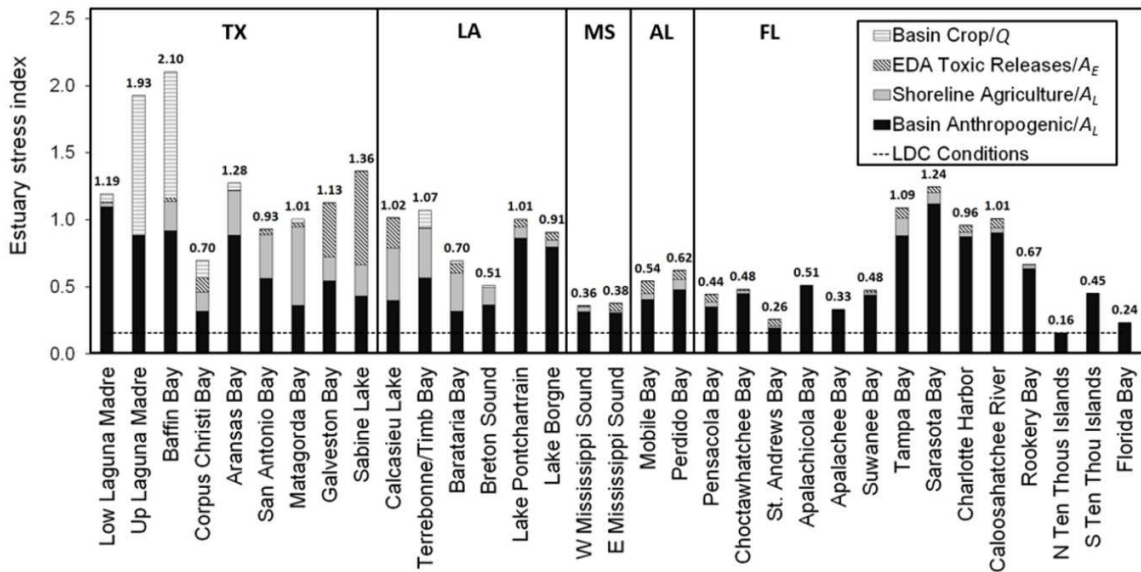
440

441 *3.3 Assessment of Estuary Level Anthropogenic Stress*

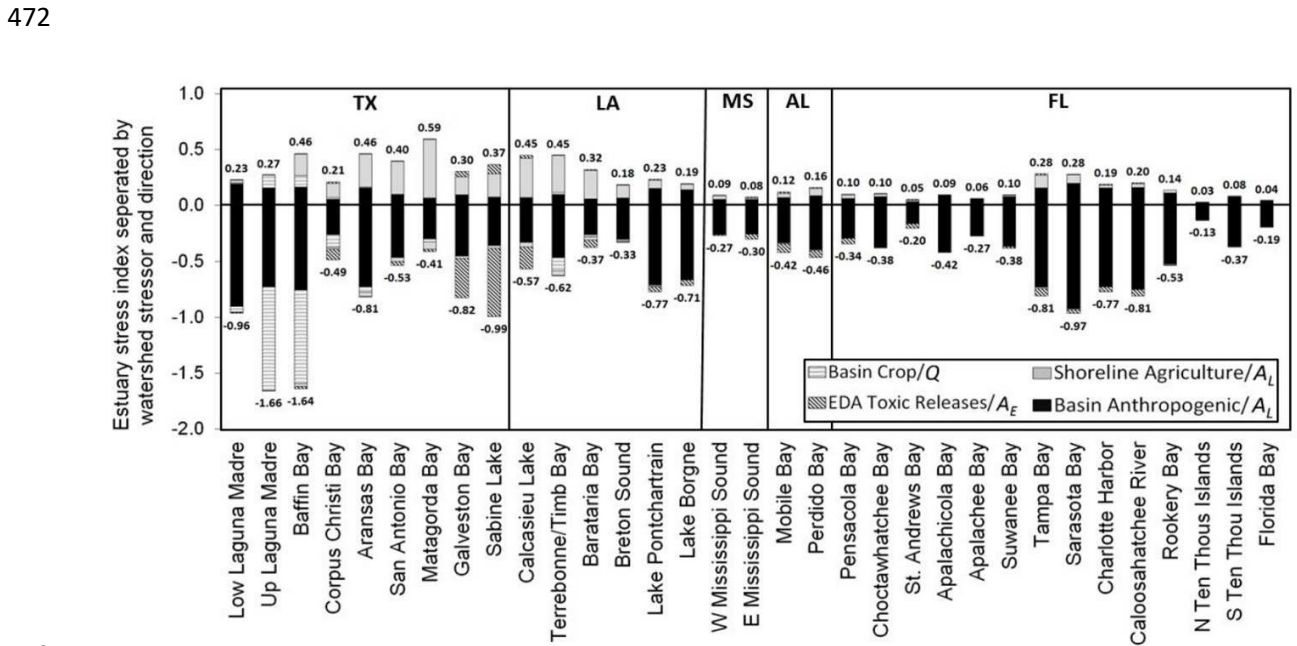
442 The aggregated intensity of anthropogenic stress for each GoM estuary was combined to form an
 443 estuary stress index based on the mean absolute change in log odds of species presence,
 444 comparing current to benchmark (LDC or MDC) conditions (Fig. 2). We used the mean absolute
 445 value of both positive and negative coefficients, reflecting that increases in stress-tolerant species
 446 can indicate estuary disturbance in the same way that declining sensitive species do (Felley 1987;
 447 Caddy 2000; de Leiva Moreno et al. 2000; Chesney and Baltz 2001; Lewis et al. 2011). The
 448 estuary stress index provides a concise and quantitative way to estimate the amount of
 449 anthropogenic stress affecting each estuary, and since values are additive in log odds, the
 450 contributions of individual stressors can be observed separately (Fig. 2). Basin Anthropogenic/ A_L
 451 had the largest effect on species presence in the GoM, averaging nearly 70% of total deviations,
 452 and was most prevalent in the highly urbanized southwest FL estuaries (Sarasota Bay, Charlotte
 453 Harbor, Caloosahatchee River, and Tampa Bay) along with several estuaries in the western GoM
 454 (Baffin Bay, Upper and Lower Laguna Madre, and Aransas Bay, TX and Lake Borgne and Lake
 455 Pontchartrain, LA). Basin Crop/ Q and EDA toxic releases/ A_E both had large effects in relatively
 456 few estuaries where those stressors had elevated values (Upper Laguna Madre, Baffin Bay,
 457 Galveston Bay, and Sabine Lake, TX and Calcasieu Lake, LA) while Shoreline Agriculture/ A_L
 458 was prevalent in several TX and LA estuaries.

459 Anthropogenic influence was also split between positive and negative effects (Fig. 3) in order to
 460 assess in which direction watershed stressors were impacting GoM estuaries. Both positive and
 461 negative components exist for the same stressor because some of the 57 species modeled are
 462 positively related to the stressor while other species are negatively related to the same stressor
 463 (Table 6; SM 4). The magnitude of negative impacts are significantly larger for Basin
 464 Anthropogenic/ A_L , Basin Crop/ Q , and EDA toxic releases/ A_E , while the opposite is true for
 465 Shoreline Agriculture/ A_L (Fig. 3).

466



467
 468 **Fig. 2** Estuary stress index is the mean anthropogenic effect in logs odds based on the absolute
 469 value of model coefficients. The x-axis represents minimally disturbed conditions (MDC; all
 470 anthropogenic stressors = 0) and the dotted line represents least disturbed conditions (LDC) for
 471 the GoM (SM 1).



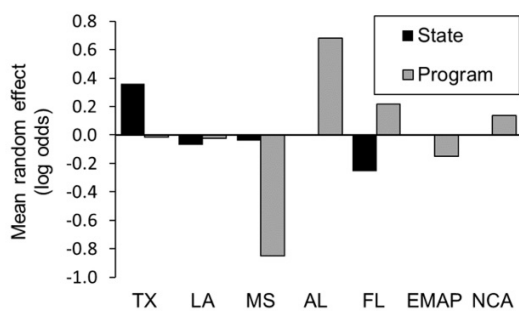
473
 474 **Fig. 3** Mean anthropogenic effect by estuary separated by positive and negative effects. The sum
 475 of the mean positive and mean negative effects equal the total absolute effect shown in Figure 2.

477 3.4 Indicator Species

478 Modeling individual species allowed for the identification of fish and invertebrate species whose
479 presence had significant negative or positive relationships with anthropogenic stressors. Species
480 with the largest negative coefficients for Basin Anthropogenic/ A_L were Atlantic Moonfish
481 (*Selene setapinnis*), Gulf Butterfish (*Peprilus burti*), Bay Whiff (*Citharichthys spilopterus*),
482 White Shrimp (*Litopenaeus setiferus*), Blue Catfish (*Ictalurus furcatus*), Bluntnose Jack
483 (*Hemicaranx amblyrhynchus*), Sea Bass (*Centropristis philadelphica*) and Fringed Flounder
484 (*Etropus crossotus*) (SM 4). Species with large negative relationships with EDA toxic
485 releases/ A_E were Gulf and Chain Pipefish (*Syngnathus scovelli*, *Syngnathus louisianae*), Pink
486 Shrimp (*Farfantepenaeus duorarum*), Atlantic Threadfin Herring (*Opisthonema oglinum*),
487 Atlantic Stingray (*Dasyatis sabina*), Blackcheek Tonguefish (*Symphurus plagiusa*), Silver Perch,
488 and Blue Crab (*Callinectes sapidus*). Several species had consistently positive responses to
489 anthropogenic stressors including: Black Drum (*Pogonias cromis*), Gafftopsail Catfish (*Bagre
490 marinus*), Hardhead Catfish (*Ariopsis felis*), White and Striped Mullet (*Mugil curema*, *Mugil
491 cephalus*), and Lined Sole (*Achirus lineatus*).

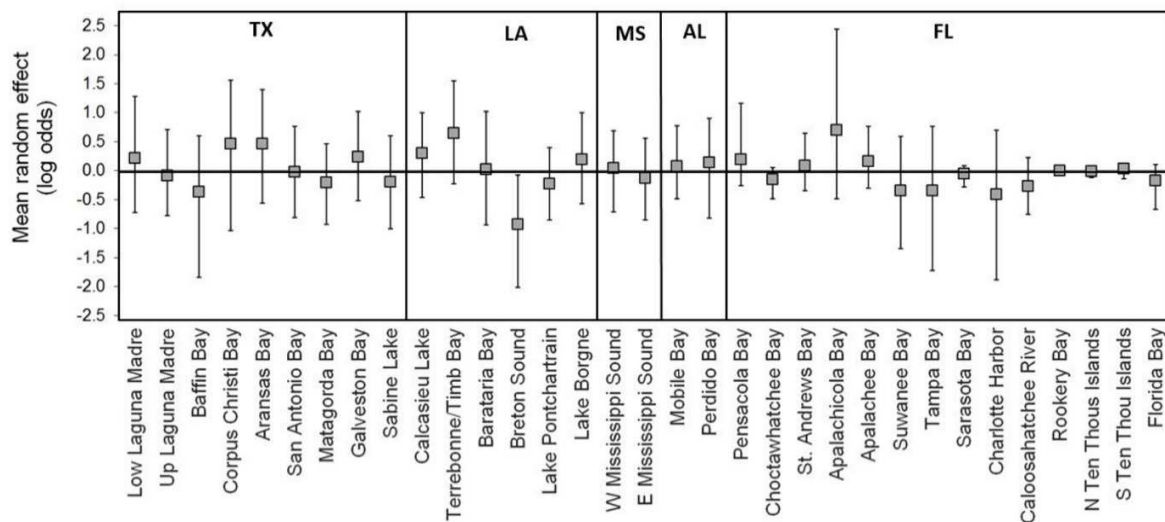
492 3.5 Random Effects and Seasonality

493 State random effects were included to account for the potential impact of state-level fishing
494 regulations on species presence, as well as large-scale natural variations in habitat that may occur
495 at the state level. In addition, program random effects were included to account for variation
496 among different trawl programs (efficiency and effort). State and program random effects were
497 averaged across all species (Fig. 4). Mean state effects ranged from -0.3 (FL) to 0.4 (TX) and
498 program effects vary from -0.8 (MS) to 0.6 (AL) suggesting substantial variability in the effects
499 of different program sampling methods on the probability of species presence in trawl events.
500 Seasonal effects were included as categorical variables. Fall had the highest mean effect while
501 mean summer, spring, and winter offsets were 0.38, 0.36, and 0.38 lower (on log odds scale),
502 respectively. It is important to note that event-level temperature and salinity measurements were
503 also included as predictors in the model, so that these seasonal effects represent variability that
504 exists in addition to seasonal patterns in temperature and salinity.



505
506 **Fig. 4** Mean state and program random effects
507 across all fish and invertebrate species.
508

509 The magnitudes of estuary-level random effects were determined by averaging across all species
 510 (Fig. 5). The relatively high random effects observed in estuaries such as Aransas Bay, San
 511 Antonio Bay, TX, Terrebonne/Timbalier Bays, LA, and Apalachicola Bay, FL point to estuaries
 512 where species presence was higher than would be expected based on the predictors and other
 513 random effects included in the models. Similarly, estuaries such as Matagorda Bay, TX and
 514 Breton Sound, LA had a lower probability of presence than would be expected. In such estuaries,
 515 other biophysical or watershed factors that were unaccounted for by this analysis may be
 516 affecting habitat quality.



517
 518 **Fig. 5** Estuary random effects. Grey square indicates the mean (across species) and error bars
 519 show the 0.1 and 0.9 quantiles. Positive random effects imply species presence was under
 520 predicted by the other components of the model. Negative random effects imply the opposite.
 521 Rookery Bay does not have an estuary random effect because it had no trawl samples.

522
 523 **4. Discussion**

524 *4.1 Anthropogenic Stressors*

525 Our study contrasts with previous biological assessment efforts that sought to characterize
 526 biological condition, but which were not designed to quantitatively relate that condition to
 527 watershed anthropogenic stressors (Deegan et al., 1997; Macauley et al., 1999; Summers, 2001;
 528 Hughes et al., 2002; Meng et al., 2002; Harrison and Whitfield, 2004, 2006; Breine et al., 2007;
 529 Coates et al., 2007; Uriarte and Borja, 2009; Breine et al., 2010; Delpech et al., 2010; Jordan et
 530 al., 2010; Cabral et al., 2012). Previous work in the U.S. (e.g., Greene et al., 2015) has
 531 characterized the magnitudes of anthropogenic stressors in estuarine watersheds, but there has
 532 been limited progress toward quantitatively linking these stressors to the biological condition of
 533 estuaries. In Europe, Teichart et al. (2016) used ecology quality ratios (EQR) scores from seven
 534 European countries to determine water quality stressors that affect estuarine biological condition.

535 Though this was a step toward linking biological condition to watershed stressors, there are still
536 questions about how comparable EQR ratios are among countries (Poikane et al., 2014) and
537 basin-level stressors were not explored. In this study, we demonstrate a linkage between
538 watershed stressors at three spatial scales (basin, EDA, and shoreline) and estuarine biological
539 condition using regression models that relate watershed stressors to the log odds of species
540 presence while accounting for natural variation and other potentially confounding effects (e.g.,
541 systematic biases associated with different sampling programs). We also develop a quantitative
542 index of estuary stress using mean stressor coefficients (averaged across species) and the
543 variability in stressors across estuarine watersheds (Figs. 2, 3). We found widespread and largely
544 negative influences of landscape stressors on both fish and invertebrate species. Our results
545 suggest that estuaries east of the Mississippi delta to the Florida panhandle are, in general, less
546 impacted by watershed stress than other systems in the GoM. Our indicator screening process
547 also provides new insight into species-level responses to estuary stressors, revealing species
548 indicators on both the tolerant and sensitive sides of the ‘stress-tolerance gradient’ (Whittier et
549 al. 2007).

550 Our results indicate that human activities in upstream watersheds have statistically significant
551 and largely negative relationships on a substantial portion of estuarine fish and invertebrate
552 species. The percent of anthropogenic land cover in river basins ($\text{Basin Anthropogenic}/A_L$) has
553 the strongest relationship, affecting 22 species (17 negatively) in the multi-stressor models
554 (Table 6) including five of the ten most prevalent species in the GoM (SM 4). It accounts for
555 nearly 70% of all absolute deviations from MDC (Fig. 3). The “Anthropogenic” land category is
556 a composite of urban, crop, and pasture, and it generally outperformed the individual land
557 categories in terms of predictive performance (Table 5). Nonetheless, it is interesting to consider
558 whether one of the three land cover classes (urban, crop and pasture) plays a dominant role. The
559 high ranking of crop and agriculture in single stressor models (Table 5) coupled with the fact that
560 agriculture comprises 69% of anthropogenic land use within our study area, suggests that it is
561 perhaps the dominant form of land cover leading to estuary disturbance. To further explore this,
562 we repeated the single-stressor modeling three times, omitting estuaries in FL, TX, and the
563 central Gulf states (LA, MS, and AL), in turn, as a simple sensitivity analysis. Basin
564 $\text{Anthropogenic}/A_L$ remained the top stressor when both the central state estuaries and the TX
565 estuaries were omitted, but Basin Crop/A_L became the top stressor when FL estuaries were
566 omitted. The increased importance of Basin Crop/A_L when Florida estuaries are excluded implies
567 that crop production might be more influential in the western GoM. Of note, the categorical
568 variables representing hypoxic, toxic algal, and eutrophic condition estimates (Bricker et al.
569 2008) were not found to be significant predictors in single-stressor models, suggesting these
570 metrics, as calculated and used in this study, are not primary drivers of species presence in GoM
571 estuaries.

572 Basin Crop/Q , reflecting crop areas normalized by average flow rate, is significantly related to
573 21 species (18 negatively). This stressor is likely a proxy for the concentration of crop-related
574 pollutants entering a system. Basin Crop/Q was highest in Baffin Bay and Upper Laguna Madre,
575 two hyper-saline southern TX estuaries, and elevated in Corpus Christi Bay, TX and

576 Terrebonne/Timbalier Bays, LA as well (SM 1). These results imply that as estuary flushing
577 decreases, estuaries become more susceptible to pollution. Hydrologic alteration as well as
578 consumptive water use by urban and agricultural sources can reduce water inflow to estuaries
579 and the dilution of contaminants in surface water (Harwell, 1997; Day et al., 2000; Flannery et
580 al., 2002).

581 The number of toxic release sites in the estuarine drainage area (EDA) normalized by the estuary
582 area is significantly related to 12 species (10 negatively). Five of the negative relationships are
583 with species associated with bottom environments (e.g., Atlantic Stingray, Blue Crab, Pink
584 Shrimp, Blackcheek Tonguefish, and Hog Choker (*Trinectes maculatus*); SM 4). Galveston Bay
585 and Sabine Lake, TX and Calcasieu Lake, LA have the highest historical levels of toxic releases,
586 potentially due to upstream petrochemical plants (USEPA 2015; Blackburn 2015). Our findings
587 are consistent with previous research reporting measurable negative effects of toxic
588 contamination and releases on estuarine organisms, particularly in the benthos (Pearson and
589 Rosenberg, 1978; Boesch and Rosenberg, 1981; Malins et al., 1985; Engle et al., 1994; Macauley
590 et al., 1999; Brown et al., 2000).

591 The percent of the estuarine shoreline buffer comprised of agricultural land cover is the only
592 anthropogenic stressor identified to be predominantly associated with an increase in species
593 presence. It is significantly related to 13 species; 12 of these positively. Estuaries with shoreline
594 agriculture greater than 10% are generally found in TX and LA (SM 1). One possible
595 explanation for this finding is a positive response to incipient stress as observed in other estuary
596 studies (Jordan and Vaas, 2000; Jordan and Smith, 2005). A possible mechanism for a positive
597 response to stress could be greater productivity and diversity due to nutrient enrichment (i.e., a
598 release from oligotrophy; Nixon and Buckley, 2002), or by a combination of stressors that has
599 not reached a threshold for causing negative effects (Jordan et al., 2010). Alternatively, long-
600 term effects of nutrient enrichment (and overfishing) have been shown to lead to the dominance
601 of pelagic over benthic/demersal species (Caddy, 2000; de Leiva Moreno et al., 2000). Nine of
602 the 12 species positively linked to shoreline percent agriculture are pelagic species (e.g., Gulf
603 Menhaden, Atlantic Bumper (*Chloroscombrus chrysurus*), Ground Mullet (*Menticirrhus*
604 *americanus*), and White Mullet; SM 4) and two more are catfish species which thrive in high
605 nutrient waters (*Ariopsis felis*, *Bagre marinus*). More investigation into the positive relationship
606 between shoreline agriculture and the presence of several species is warranted.

607

608 4.2 Assessment of Estuary Level Anthropogenic Stress

609 We adopted both MDC and LDC as possible benchmarks against which to compare estuary
610 disturbance in order to demonstrate how our index of estuary stress could be used with both;
611 however, each of these potential benchmarks have limitations. MDC requires a somewhat greater
612 degree of model extrapolation, while LDC is not referenced to an absolute ecological state
613 making it susceptible to shifting baselines as watersheds continue to develop over time (Pauly,
614 1995). LDC conditions, as defined in this study, do not deviate greatly from MDC (0.16 change
615 in log odds; Fig. 2). LDC for both Shoreline Agriculture/ A_L and EDA toxic releases/ A_E were zero

616 (i.e., equal to MDC) while LDC for Basin Crop/ Q is essentially the same as MDC. The minimum
617 LDC value for Basin Anthropogenic/ A_L was 8.9% and this stressor explains the majority of the
618 difference between MDC and LDC. The application of LDC across such a large system as the
619 GoM might be debatable (Stoddard et al., 2006); however, comparing the MDC and LDC
620 benchmarks in Fig. 2 suggest the overall disturbance pattern indicated by our results should be
621 fairly robust to small changes in benchmark assumptions. Estuary rankings essentially remain the
622 same whether they are based on deviation from LDC or MDC conditions since both benchmarks
623 are constant across the GoM and would only change if sub-regional benchmarks are developed
624 on the assumption that baseline conditions vary by region. Another approach that might lead to
625 different estuary ranks would be if the estuary stress index was defined as only the negative (i.e.,
626 negative component in Fig. 3) or the net effect of watershed level stressors (i.e., the difference
627 between the positive and negative components in Fig. 3) for specific subsets of species of
628 interest. If system productivity (i.e., more fish for anglers) is of interest, then an estuary stress
629 index could be created specifically for species of interest for commercial and recreational
630 fisherman. However, both of these alternatives ignore the value of stress-tolerant species as
631 potential indicators of anthropogenic influence (see below; Whittier et al., 2007).

632 *4.3 Indicator Species*

633 The estuary stress index proposed here is based on the responses of widely distributed fish and
634 invertebrate species to different landscape anthropogenic stressors that provide insights into
635 important regional indicator species. Forty-eight of the 57 species studied are significantly
636 related to at least one anthropogenic stressor in the multi-stressor models (SM 4). While the
637 overwhelming direction of response is negative, responses of species with apparent stress
638 tolerance (i.e., those that responded positively to increasing stressor levels) are also present (see
639 Section 3.4). Tolerant species tend to have greater behavioral and diet plasticity, greater niche
640 breadth and may benefit from the absence of more sensitive competitors (Vázquez and
641 Simberloff, 2002; Swihart et al., 2003; Devictor et al., 2008; Wilson et al., 2008; Segurado et al.,
642 2011).

643 *4.4 Hierarchical Modeling*

644 The ability to model variability at multiple scales (i.e., estuaries, states, programs) is an
645 important advantage of hierarchical modeling in this application. Critically, it allowed us to
646 perform statistically valid hypothesis testing concerning which watershed anthropogenic
647 stressors are significantly related to species presence. The hierarchical approach can be
648 contrasted with more traditional “no pooling” and “complete pooling” approaches to regression
649 modeling (Gelman and Hill, 2006; Qian et al., 2010; Cuffney et al., 2011). If we had developed
650 independent models (or parameter estimates) for each estuary (no pooling), then it would be
651 infeasible to statistically assess the significance of stressors acting across multiple estuaries. On
652 the other hand, if all data were combined into a single model, but random effects were omitted,
653 then we would be treating each trawl sample as statistically independent, failing to address the
654 substantial intraclass correlation that exists among samples from the same estuary, state, and
655 program (Gelman et al., 2014). As an example, single-stressor models (Eq. 2) were run omitting

656 the estuary random effect (α_{estuary}) and 40 of the 47 predictor variables listed in Table 4 were
657 significant for over 80% of species; an implausible result related to the failure to account for
658 intraclass correlation.

659 In addition, random effects allow us to characterize variability over different scales (i.e., sample,
660 estuary, state, and programs) (Wikle, 2003a; Cressie et al., 2009). For example, program-level
661 random effects allow us to assess the variability associated with differences in sampling
662 protocols. The MS program only recorded 23 of the 57 modeled species, resulting in a much
663 lower mean program random effect when compared to other state and federal programs (Fig. 4).
664 State random effects are also of interest, with TX having the highest state random effect and FL
665 having the lowest, though the variation across states was less than the variation across programs
666 (Fig. 4). Exploring the causes of state-level variation could be a subject for future research, as
667 this variation could plausibly be related to either state fishing regulations, biogeography, or other
668 factors that vary over a large spatial scale. The ability to discern between state and program level
669 random effects is critical, as state random effects reflect actual differences in species prevalence,
670 whereas program random effects reflect sampling efficiency. Distinguishing between state and
671 program random effects within the model was only possible because of the two federal interstate
672 programs (EMAP and NCA), and this is an important reason to continue these federal trawl
673 programs in the future.

674 *4.5 Summary and Future Directions*

675 This study identifies watershed anthropogenic stressors that are most strongly associated with
676 fish and invertebrate species presence, and then used the modeled stressor relationships to
677 develop an index of watershed stress for estuaries across the GoM. The hierarchical generalized
678 linear model used here provides a tool to control for variability that occurs among programs,
679 estuaries, and states, as well as among individual trawl samples. A large percentage of GoM
680 species exhibited negative associations with anthropogenic stressors, but some positive
681 associations were present as well. The percent anthropogenic land cover (urban, crop, and
682 pasture) within an estuary's drainage basin was consistently ranked as a leading predictor of
683 estuary disturbance. When determining deviations from benchmark conditions (Fig. 2), we
684 implicitly assumed that the statistically significant predictor variables identified through
685 hierarchical modeling are, in fact, drivers of species presence/absence. However, the empirical
686 relationships developed in the single and multi-stressor models presented here do not directly
687 demonstrate causal relationships, as it is well known that correlation is not necessarily causation.
688 The results do, however, provide a data-supported line of evidence relevant to large-scale
689 estuarine assessments and fisheries management in the GoM. Past studies have found negative
690 effects of nitrogen loading, eutrophication, and urbanization on estuary species (Dauer et al.,
691 2000, Diaz and Rosenberg, 2008; Coll et al., 2010), but our results are perhaps the most
692 compelling evidence to date for widespread population-level effects on fish and invertebrate
693 habitats in GoM estuaries by watershed stressors, given the comprehensiveness of the datasets
694 used and the spatial extent of the study. We expect this study to motivate and focus future
695 research efforts to understand the mechanisms by which watershed stressors affect estuary

696 habitats. Our findings may also be useful to help guide and prioritize watershed restoration
697 activities.

698 While the current study focuses on species-level responses, other studies have focused on the
699 responses of biological assemblages (i.e., metrics of community composition, structure, and
700 function) to regional stressor gradients (Lewis et al., 2007; Piazza and La Peyre, 2009).
701 Acknowledging the potential advantages to species-specific indicators (see Introduction), our
702 approach could be extended to consider the presence/absence of species functional groups that
703 share common habitat, life history, or feeding strategies. Modeling functional groups may help
704 corroborate the patterns documented here and provide further insights into the mechanisms by
705 which landscape stressors act on biological communities. Further, modeling functional groups
706 could facilitate comparison of results to previous studies that have relied on community metrics
707 for estuarine biological assessment (Engle et al., 1994; Summers, 2001; Hughes et al., 2002;
708 Jenkins, 2004; Jordan et al., 2010).

709 This hierarchical approach could be extended to consider species abundance, which is related
710 (imperfectly) to the number of individuals of a given species collected in a trawl event. An
711 assessment of how species abundance responds to anthropogenic stressors could also be
712 valuable, particularly for species of social or economic importance. Many of the species found to
713 have sensitivities to stressors in this study are also targeted by the regional commercial fishery
714 (including shrimps, Gulf Menhaden, and Blue Crab), and species that are heavily targeted in
715 recreational fisheries (Spotted Seatrout, Sand Seatrout (*Cynoscion arenarius*), Silver Seatrout
716 (*Cynoscion nothus*), Ground Mullet, and Atlantic Croaker (*Micropogonias undulates*); NMFS,
717 2014). A natural extension of this research is to focus on the responses of economically valuable
718 species in terms of both their probabilities of occurrence and abundances.

719 Finally, the hierarchical modeling approach developed in this study could be enhanced to
720 consider additional spatial and temporal variability. Our approach does consider event-level
721 variables like temperature and salinity that explain some intra-estuary variability, but which
722 cannot be directly related to stressors. Like other studies before ours (Jordan et al., 2010), our
723 assessment is not capable of estimating variation in biological condition at a sub-estuary scale. If
724 trawl samples could be organized into different estuarine subsections (perhaps based on estuary
725 geomorphology), then species presence could be related to more localized stressors associated
726 with shoreline development. Shoreline hardening associated with sea walls has been shown to
727 reduce both richness and abundance of species in near shore areas (Peterson and Lowe 2009;
728 Gittman et al. 2016; Dethier et al. 2016), but precise estuary-level estimates of shoreline
729 hardening were not available for this study (Gittman et al. 2015). Regarding temporal variability,
730 future analyses could explore how species presence changes over time, with alterations in
731 watershed development and basin flows as potential predictor variables. A better understanding
732 of temporal trends in estuary condition could further prioritize watershed management and
733 restoration activities.

734

735

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