



Changes in arthropod communities as black mangroves *Avicennia germinans* expand into Gulf of Mexico salt marshes

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Received: 19 March 2018 / Accepted: 25 September 2018 / Published online: 5 October 2018
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Abstract

Climate change is driving poleward shifts in species distributions worldwide. In the Gulf of Mexico (GOM), warming temperatures foster black mangrove (*Avicennia germinans* L.) expansion into GOM wetlands replacing wetland plants including *Spartina alterniflora* Loisel, *Salicornia depressa* L., and *Batis maritima* L. We investigated insect community assemblages in wetlands with and without *A. germinans* to assess potential effects of *A. germinans* expansion on insect fauna. Insect abundance, biomass, richness, diversity, community structure, and feeding guild composition were measured in both the spring and the fall across three levels of *A. germinans* abundance. Insect abundance and biomass were larger in both the spring and the fall in wetlands where *A. germinans* abundance was low. Significant differences in community structure were associated with the presence of *A. germinans*. Feeding guild composition was also different in wetlands containing *A. germinans*, having less predator biomass. Shifting vegetation caused by climate change can alter insect communities in coastal wetlands, illustrating the need for a more comprehensive understanding of climate change effects on fauna in response to shifting foundation plant species.

Keywords Climate change · Vegetation shift · Insect · Community change

Introduction

Coastal ecosystems are experiencing significant effects of climate change (e.g., Loarie et al. 2009), which is of concern because two-thirds of the planet's human population and some of its most productive ecosystems and biodiversity hot spots occur within these areas (Agardy et al. 2005). Challenges associated with climate change are and will continue to affect coastal ecosystems through sea level rise, ocean acidification as well as warming temperatures (e.g., Micheli et al. 2008; Craft et al. 2009; Koch et al. 2012; Armitage et al. 2015). Warming temperatures are associated with distribution changes in many taxa (Thomas et al. 2004), and they are driving a poleward expansion of species across the globe (Hickling et al. 2006). Tropical species are becoming more abundant in temperate areas, which has significant effects on biodiversity and ecosystem function (Hickling

et al. 2006; Micheli et al. 2008; Vergés et al. 2014; Guo et al. 2017). For example, tropical herbivores may move into temperature regions and alter occurrence and distribution of aquatic vegetation (Vergés et al. 2014). Warming temperatures are also facilitating shifts in foundation species, which may fundamentally alter community structure and function as well as ecosystem processes (Ellison et al. 2005, Micheli et al. 2008; Armitage et al. 2015; Guo et al. 2013).

Plants are the primary source of food and habitat for consumers, and a shift in the abundances or distributions of existing plant fauna may alter the composition of and interactions between the organisms that inhabit them (Gratton and Denno 2005, 2006; Armitage et al. 2015). For example, invasion by the common reed *Phragmites australis* (Cav.) changed salt marsh faunal composition on the US Atlantic Coast (Osgood et al. 2003; Kimball et al. 2010). In estuaries, warming temperatures have facilitated seagrass species shifts in North Carolina and Texas, which were associated with changes in faunal assemblages and primary productivity (Micheli et al. 2008). In Gulf of Mexico (GOM) wetlands, milder winters without severe freezing events have allowed black mangroves *Avicennia germinans* L. to expand northward into salt marshes, displacing native wetland plants

Handling Editor: Miriama Malcicka.

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(Osland et al. 2013; Cavanaugh et al. 2014). This change is associated with significant changes in microclimates, soil organic content, sedimentation rates, and associated birds (Guo et al. 2017). Salt marshes provide numerous ecosystem services including storm surge and coastal erosion protection, carbon sequestration, primary production, and habitat for numerous aquatic and terrestrial species (Pennings and Bertness 2001). Alteration of this vital ecosystem through changes in vegetation make-up could dramatically alter coastal wetland food webs, change ecosystem properties, and create new niches for invasive species (Gedan et al. 2009).

The salt marsh–mangrove barrier exists at or near *A. germinans* temperature thresholds (Record et al. 2013), suggesting that even a minimal increase in mean annual temperature accompanied by a decrease in severe freezing events (colder than $-4\text{ }^{\circ}\text{C}$) could lead to extensive increases in *A. germinans* distribution, altering the structure and function of coastal wetlands (Cavanaugh et al. 2014). Cold temperatures have historically limited the northern range limit for *A. germinans* to around 30°N (Kangas and Lugo 1990). Over the last few decades, a decrease in the frequency of sustained severe freezing events has allowed the area of mangrove forests to double at the northern end of this range in the Gulf Coast of the United States (Cavanaugh et al. 2014). In marshes near Corpus Christi, TX, USA, mangrove cover has expanded from less than 100 acres to more than 27,000 acres since 1980 (Montagna et al. 2011.; Armitage et al. 2015). *A. germinans* encroachment alters below ground properties (e.g., soil characteristics, biomass, carbon storage), and these effects are more intense in drier wetlands like those in the Western GOM (Yando et al. 2016).

In some areas, *A. germinans* dominates the lower tidal elevations (Geldenhuys et al. 2016), but in the Western GOM, *A. germinans* tends to occupy the higher tidal elevations replacing upper elevations of *Spartina alterniflora* Loisel and other marsh plants in the process (Smee et al. 2017). *S. alterniflora* remains at the lowest tidal elevations in a small border surrounding a dwarf *A. germinans* forest, and *S. alterniflora* facilitates *A. germinans* seedlings by creating a buffer against cold temperatures (Guo et al. 2013) and by trapping the *A. germinans* propagules from becoming dispersed outside of optimal retention ranges (Peterson and Bell 2012). *A. germinans* are replacing the existing flora in the Southern GOM, and anticipated warming trends are likely to promote *A. germinans* expansion poleward, causing further displacement of marsh plants.

Insects are ubiquitous in coastal wetlands (Pennings and Bertness 2001). Insect families are generally restricted in diet to a small group of related plants due to coevolutionary defense strategies (Futuyma and Mitter 1996; Burrows 2003; Nagelkerken et al. 2008). For instance, tropical plants generally invest heavily in chemical defenses relative to temperate plants (Coley 1998). Evidence of coevolution has been seen

in arthropods (Coley and Aide 1990), fish (Bertness et al. 1981), crustaceans (Heck and Wilson 1987), and various intertidal species (Menge and Lubchenco 1981). In general, *A. germinans* invests more in defense strategies to combat insect herbivory than do temperate marsh plants (He and Silliman 2015). Thus, replacement of a temperate plant species by a tropical one such as *A. germinans* might affect grazing rates and energy transfer. Insects play an important role in decomposition and the cycling of nutrients through ecosystems as well as being a major food source for fish, birds, amphibians, reptiles, and other invertebrates making them essential for habitat functional integrity (Angermeier and Karr 1994). Aquatic fauna were significantly different in wetlands dominated by *A. germinans* (Diskin and Smee 2017; Smee et al. 2017), but the effects of mangrove expansion on salt marsh insect communities remain largely unexplored in the GOM.

Predicting potential effects of *A. germinans* expansion on insects is difficult. By displacing other plants and developing a monoculture, *A. germinans* may lower insect diversity and abundance. Alternatively, by creating a new ecotone, *A. germinans* might increase insect diversity and enhance wetland food webs, particularly in wetlands transitioning from domination by grasses and succulents to dwarf *A. germinans* forests. The purpose of this study was to determine what changes, if any, are occurring in the GOM estuarine wetland insect communities as *A. germinans* replace salt marshes and attempt to identify patterns associated with this change.

Materials and methods

Study sites

Samples were collected from marshes in estuaries near Port Aransas and Rockport, TX, USA. *A. germinans* has become well established in many areas but is nearly completely absent in others. In some places, *A. germinans* has displaced all marsh plants common in higher intertidal elevations, such as *Salicornia depressa* L. and *Batis maritima* L., as well as the upper tidal elevations typically occupied by *S. alterniflora* (Smee et al. 2017). In these areas with established and abundant *A. germinans*, *A. germinans* has formed dense monocultures at higher intertidal elevations excluding all other plants and has restricted *S. alterniflora* to narrow bands at the lowest tidal elevations (Fig. 1). Thus, marshes change from a diverse plant assemblage containing grasses and succulent plants to one dominated by *A. germinans*. We categorized marshes by the abundance of *A. germinans*, which was inversely related to the abundance of *S. depressa*, *B. maritima*, and *S. alterniflora*. Abundance was based on measurements of vegetation width (distance from lowest to highest tidal elevations, Diskin 2016). For this



Fig. 1 Wetlands with abundant (top) black mangroves *Avicennia germinans*. Author Loveless and ETEAMS intern Erin Urban collect insects in a wetland with few *A. germinans* (bottom)

study, marshes were grouped into three types based on the relative abundances of these plants: (1) rare *A. germinans*, and *S. depressa*, *B. maritima*, *S. alterniflora* abundant, (2) intermediate *A. germinans* co-occurring with similar abundance of *S. alterniflora*, *S. depressa*, and *B. maritima*, and (3) abundant *A. germinans*, in which other plants were in low abundance.

Sampling design

Insects were collected within each wetland type in June (late spring) and October (fall) of 2016. *A. germinans* were flowering during spring sampling and not during the fall with no other vegetation flowering while samples were taken. Eight samples were collected from each wetland type (24 from each season, 48 total). Insect specimens were collected using a suction sampler made from a converted leaf blower-vac (Buffington and Redak 1998). Samples were taken using a sweeping motion from side to side in a near 180° arc with 1 sweep made, then 1 step forward taken, and another sweep

made, until 15 sweeps were completed and 15 constituted one sample (Buffington and Redak 1998). Catch per unit effort (CPUE) is therefore defined in this paper as the number of individuals per 15 sweeps of the suction sampler. Other techniques such as sweep nets and light traps were tested and determined to be less effective in capturing insects (Loveless 2017). We collected fewer insect species and overall many fewer individuals using these methods as compared to the suction sampler. Wetland plants were also searched by hand to ensure that insects had not been missed with the vacuum sampler. Insects such as lepidopterans that are not effectively sampled with a suction sampler were not found (Loveless 2017).

Once collected, insects were stored in 95% ethanol and brought to the lab for sorting, identification, and enumeration. The primary focus of this study was on insect taxa; however spiders were enumerated and simply identified as arachnid for both order and family. Feeding guilds were grouped using the classifications of Papp (2002) and Sinu and Sharma (2013): fungivore, herbivore, parasite, parasitoid, predator, and saprophage. Herbivores were further separated into chewers, sap feeders, and a third miscellaneous category (Pomeroy and Wiegert 1981) encompassing all remaining groups (e.g., gall formers, nectar feeders). Blood-sucking insect families (e.g., Culicidae, Tabanidae) were considered parasites, and detritivores were characterized as saprophagous.

Data analysis

Insect communities were compared among seasons (spring and fall) and wetland types (*A. germinans* rare, *A. germinans* intermediate, *A. germinans* abundant) using a 2-way PERMANOVA (PRIMER™). Many more insects were collected in spring than fall; therefore, insect community differences between wetland types were compared separately among seasons. Non-metric MDS plots were created for both the spring and the fall to visualize differences among insect communities by wetland type. An analysis of similarity (ANOSIM) in PRIMER™ was performed to examine the differences in communities among wetland types in spring and fall, and a similarity percentage (SIMPER) analysis in PRIMER™ was used to determine which insect families contributed most to insect community differences among wetland types.

Univariate analyses (insect abundance, biomass, richness), and Shannon–Weiner diversity, were compared in the spring and the fall using ANOVA in JMP Pro 13.1 with wetland type as a fixed factor. Tukey post hoc tests were performed to compare pairwise differences among wetland types.

Comparisons of feeding guilds were simplified by separating wetland types into two groups: with

(intermediate + abundant) and without (rare) *A. germinans*. Many of the insects collected were small phorid flies (< 2 mm, < 2 mg) which contributed substantially to the number of individuals collected; however, they contributed very little to total biomass. Because of this, biomass contributions were used when comparing the feeding guilds between the wetland types.

Results

Community structure

Insect communities were significantly different among wetland types (Pseudo $F = 13.73_{1,42}$, $p < 0.001$) and seasons (Pseudo $F = 13.73_{1,42}$, $p < 0.001$). The interaction between wetland type and season was also significant (Pseudo $F = 8.20_{1,42}$, $p < 0.001$). MDS plots for the spring (Fig. 2a) and fall (Fig. 2b) showed distinct grouping patterns based on wetland type. Consistent with MDS, ANOSIM indicated a significant effect of wetland type in both spring ($R = 0.92$; $p = 0.001$) and fall ($R = 0.42$; $p = 0.001$).

SIMPER analysis revealed that regardless of season, insect communities were most dissimilar when wetlands with rare *A. germinans* were compared to those containing *A. germinans*, regardless of whether *A. germinans* abundance was intermediate or if *A. germinans* was established and had excluded most other plants (Tables 1, 2).

Abundance and biomass

In spring, 2947 insects from 31 families were collected. Individuals in the family Phoridae were the most abundant group collected (1136). Insect abundances were significantly different among wetland types, ($F_{2,21} = 8.24$, $p < 0.001$, Fig. 3); wetlands with rare *A. germinans* had more individuals than areas with intermediate or abundant *A. germinans*. Similarly, biomass was significantly different among wetland types ($F_{2,21} = 19.95$, $p < 0.001$, Fig. 3) and was highest in areas with rare *A. germinans*. Biomass was not different between areas with intermediate or abundant *A. germinans*.

Fewer insects were collected overall in fall: 992 individuals, 25 families. Ephydriidae was the most common family with 356 individuals. Like spring, insect abundance in the fall between wetland types was also significantly different ($F_{2,21} = 28.7$, $p < 0.001$, Fig. 3), and areas without *A. germinans* had more insects than areas with intermediate and abundant levels of *A. germinans*. Biomass was also significantly different among wetland types ($F_{2,21} = 19.95$, $p < 0.001$, Fig. 3), and as with abundance, wetlands without *A. germinans* had higher insect abundance than those containing *A. germinans* at intermediate and abundant levels. Arachnids contributed most of the fall

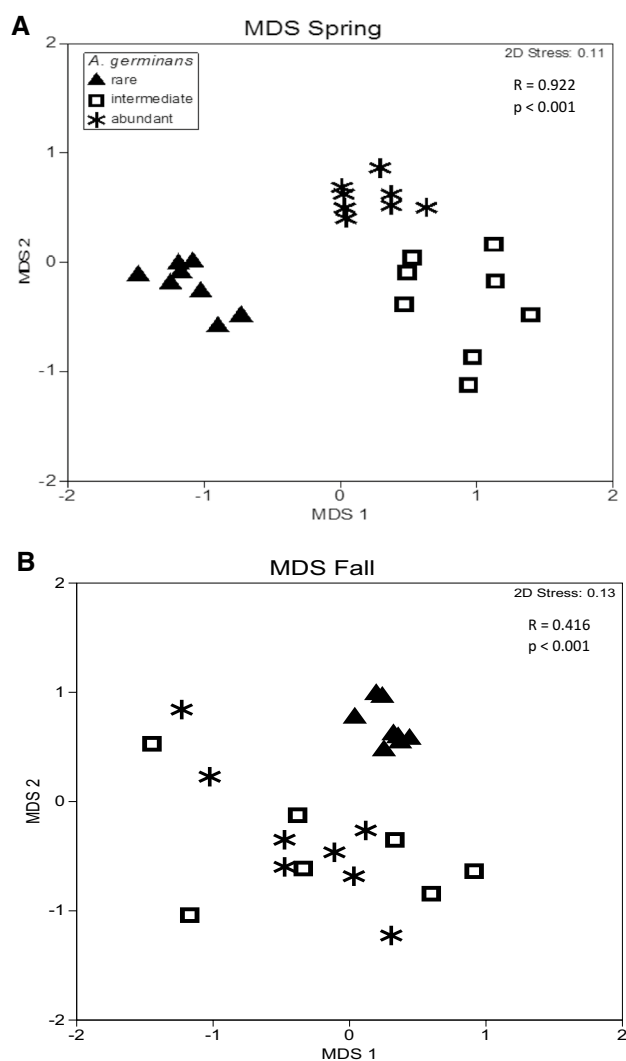


Fig. 2 nMDS plots of insect communities in wetlands with three levels of *A. germinans* presence: rare, intermediate, and abundant in the spring and fall. Global R and p values from ANOSIM are presented

biomass, comprising 3.38 g of the 10.06 g collected. Insect abundance was significantly higher in wetlands without *A. germinans* in both spring ($F_{2,21} = 13.8$, $p < 0.001$) and fall ($F_{2,21} = 38.8$, $p < 0.001$), as compared to areas with *A. germinans* (Fig. 3).

Richness and diversity

Species richness was significantly different among wetland types in the spring ($F_{2,21} = 18.8$, $p < 0.001$) and fall ($F_{2,21} = 46.5$, $p < 0.001$, Fig. 4). Richness was significantly higher in wetlands with abundant *A. germinans*. Shannon–Weiner diversity was significantly greater in wetlands containing *A. germinans* in spring ($F_{2,21} = 7.28$, $p < 0.01$) and fall ($F_{2,21} = 6.2$, $p < 0.001$, Fig. 4).

Table 1 SIMPER analysis of dissimilarities of most common organisms collected in each wetland type during spring

Organism	Wetland type (mean insect CPUE)		Rare vs. intermediate <i>A. germinans</i>	
	Rare <i>A. germinans</i>	Intermediate <i>A. germinans</i>	Dissimilarity	% Contributed
Average dissimilarity = 66.29				
Phoridae	11.06	0.92	3.23	26.74
Delphacidae	5.28	0.25	3.24	13.14
Ulidiidae	3.37	0.47	1.96	7.80
Miridae	3.17	0.72	1.7	7.06
Ephydriidae	1.42	3.54	1.74	5.58
Culicidae	1.68	0	2.78	4.44
Dolichopodidae	2.10	2.96	1.52	4.37
Cicadellidae	1.16	2.0	1.23	4.24
Organism	Wetland type (mean insect CPUE)		Rare vs. abundant <i>A. germinans</i>	
	Rare <i>A. germinans</i>	Abundant <i>A. germinans</i>	Dissimilarity	% Contributed
Average dissimilarity = 58.07				
Phoridae	11.06	1.82	13.2	22.73
Ephydriidae	1.42	6.37	7.13	12.28
Delphacidae	5.28	1.53	5.27	9.08
Dolichopodidae	0	2.53	3.65	6.28
Miridae	3.17	1.07	3.41	5.87
Ulidiidae	3.37	2.40	2.84	4.90
Sciaridae	1.71	0.25	2.15	3.70
Pieridae	0	1.41	2.06	3.55
Culicidae	1.68	0.35	2.01	3.46
Organism	Wetland type (mean insect CPUE)		Intermediate vs. abundant <i>A. germinans</i>	
	Intermediate <i>A. germinans</i>	Abundant <i>A. germinans</i>	Dissimilarity	% Contributed
Average dissimilarity = 51.51				
Ephydriidae	3.54	6.37	5.69	11.05
Dolichopodidae	0	2.53	5.04	9.79
Ulidiidae	0.47	2.40	3.96	7.68
Ceratopogonidae	2.96	1.89	3.53	6.84
Phoridae	0.92	1.82	3.34	6.49
Delphacidae	0.25	1.53	2.92	5.66
Cynipidae	1.36	0.25	2.71	5.25
Pieridae	0.38	1.41	2.46	4.77
Pyralidae	0.75	0.88	2.09	4.06
Cicadellidae	2.00	1.51	2.06	3.99
Gryllidae	1.00	1.43	1.75	3.41
Arachnid	1.39	1.82	1.73	3.37

Feeding guild composition

All eight of the defined feeding guilds were represented in each wetland type. Herbivores (chewers, sapsuckers, and miscellaneous) were generally the most abundant feeding group followed by carnivorous guilds (predators, parasitoids, and parasites, Table 3). In the fall, sites with both

intermediate and abundant levels of *A. germinans* had significantly more arachnids than other sites. Sites in which *A. germinans* were rare/absent contained an herbivore-to-carnivore ratio of 9:1 in the spring and 5:1 in the fall. In wetlands with *A. germinans* present, this ratio was 6:1 in the spring, but switched to a 1:1 herbivore-to-carnivore ratio in the fall.

Table 2 SIMPER analysis of dissimilarities of most common organisms collected in each wetland type during fall

Organism	Wetland type (mean insect CPUE)		Rare vs. intermediate <i>A. germinans</i>	
	Rare <i>A. germinans</i>	Intermediate <i>A. germinans</i>	Dissimilarity	% Contributed
Average dissimilarity = 86.07				
Ephydriidae	5.63	0.25	17.18	19.96
Cicadellidae	4.48	0	15.03	17.46
Ceratopogonidae	3.50	0.30	10.83	12.58
Miridae arachnid	3.82	0.65	10.60	12.31
Mycetophilidae	1.87	0	6.33	7.35
Phoridae	1.81	1.10	5.23	6.08

Organism	Wetland type (mean insect CPUE)		Rare vs. abundant <i>A. germinans</i>	
	Rare <i>A. germinans</i>	Abundant <i>A. germinans</i>	Dissimilarity	% Contributed
Average dissimilarity = 83.11				
Ephydriidae	5.63	0.38	16.50	19.85
Cicadellidae	4.48	0	14.78	17.79
Ceratopogonidae	3.50	0	11.62	13.98
Arachnid	3.82	0.85	9.69	11.66
Mycetophilidae	1.87	0.25	5.61	6.75

Organism	Wetland type (mean insect CPUE)		Intermediate vs. abundant <i>A. germinans</i>	
	Intermediate <i>A. germinans</i>	Abundant <i>A. germinans</i>	Dissimilarity	% Contributed
Average dissimilarity = 74.61				
Phoridae	1.10	1.11	15.71	21.06
Arachnid	0.65	0.85	12.21	16.36
Ephydriidae	0.25	0.38	7.08	9.49
Sciaridae	0.25	0.25	5.23	7.02
Lygaeidae	0.25	0.25	5.05	6.77
Ulidiidae	0.13	0.13	3.93	5.27
Ceratopogonidae	0.30	0	3.90	5.23

Fig. 3 Mean + SE insect abundance and biomass in three levels of *A. germinans* presence: rare, intermediate, and abundant in the spring and fall. Letters denote significant pairwise differences from Tukey's post hoc test

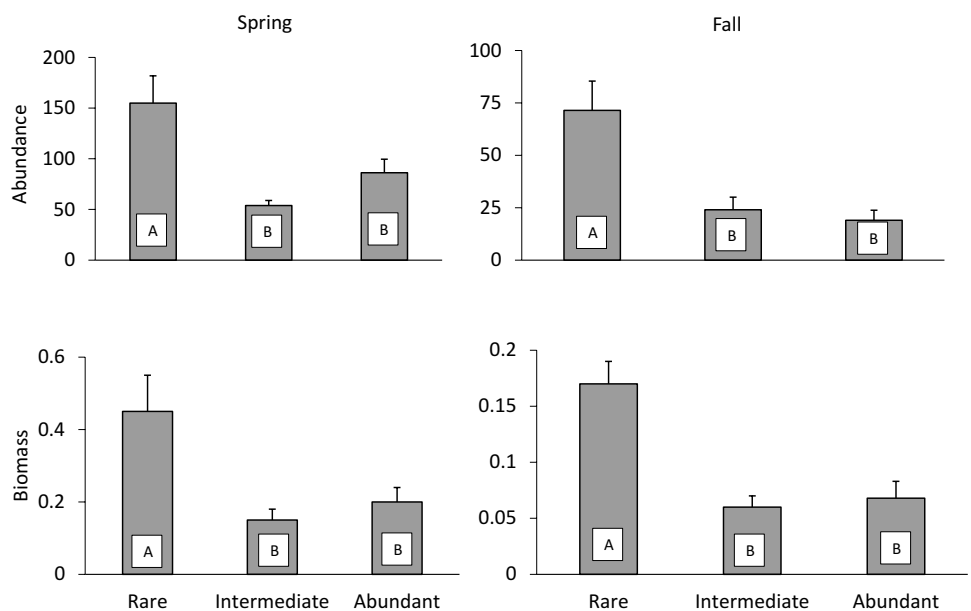


Fig. 4 Mean + SE insect species richness and Shannon diversity values in three levels of *A. germinans* presence: rare, intermediate, and abundant in the spring and fall. Letters denote significant pairwise differences from Tukey’s post hoc test

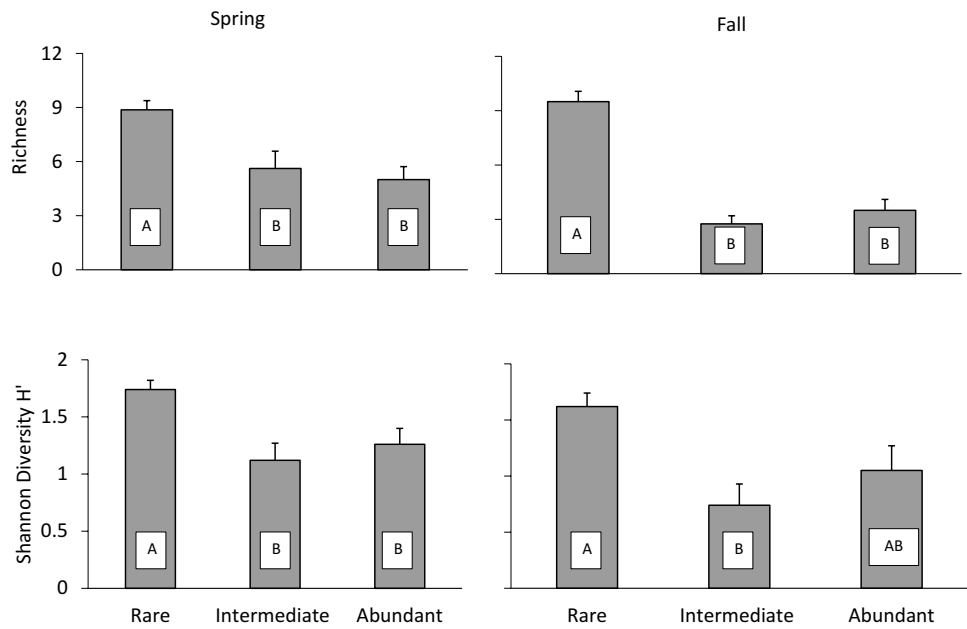


Table 3 Insect function groups by season in wetlands with and without *A. germinans*

Feeding group	<i>A. germinans</i> absent				<i>A. germinans</i> present			
	Abundance (CPUE)	% Abundance	Biomass (g)	% Biomass	Abundance (CPUE)	% Abundance	Biomass (g)	% Biomass
Spring								
Fungivore	25	1.38	0.05	0.16	15	1.34	0.032	0.16
Herbivore chewer	41	2.26	15.46	48.39	33	2.94	8.97	44.94
Herbivore misc.	134	7.39	2.083	6.52	592	52.76	5.09	25.5
Herbivore sapsucker	400	22.04	10.46	32.74	135	12.03	2.52	12.63
Parasite	71	3.91	0.33	1.03	123	10.96	0.26	1.3
Parasitoid	6	0.33	0.06	0.19	19	1.7	0.53	2.66
Predator	58	3.2	1.33	4.16	132	11.76	2.1	10.52
Saprophage	1080	59.5	2.19	6.85	73	6.51	0.34	1.7
Total	1815		31.95		1122		19.96	
Fall								
Fungivore	38	4.16	0.108	1.17	6	7.6	0.014	1.76
Herbivore chewer	11	1.2	0.18	1.94	1	1.27	0.016	2
Herbivore misc.	355	38.89	1.48	15.98	13	16.46	0.089	11.15
Herbivore sapsucker	212	23.22	3.96	42.76	10	12.66	0.114	14.29
Parasite	114	12.49	0.28	3.02	3	3.8	0.006	0.75
Parasitoid	30	3.29	0.59	6.37	6	7.6	0.09	11.28
Predator	105	11.5	2.42	26.13	19	24.05	0.43	53.88
Saprophage	48	5.26	0.25	2.7	21	26.58	0.04	5.01
Total	913		9.26		79		0.798	

Discussion

The presence of *A. germinans* was associated with reductions of other commonly occurring marsh plants in the Western GOM (Diskin and Smee 2017; Smee et al. 2017).

As climate change trends continue, it is likely that the range of mangrove forest will move northward, displacing native marsh plants. The effects of this expansion of mangrove forests on salt marsh insect communities has largely been unexplored in the GOM. Coevolution of insects and marsh plants in coastal wetlands suggests that the

disappearance of native marsh plants could significantly alter associated insect communities. Insects play an important role in nutrient cycling and energy transfer in their ecosystem; a change in insect communities could affect decomposition rates and food abundance for higher trophic levels such as fish, birds, and other invertebrates. Understanding if and how insect communities are changing as mangrove forests replace salt marsh plant species will help elucidate other potential cascading effects throughout salt marsh food webs and ecosystem functioning.

In this study, we collected and investigated insect community assemblages in wetland areas that had three levels of *A. germinans* abundance. Insects were less abundant and had lower biomass in areas with *A. germinans*. Insect communities were significantly different in areas where *A. germinans* was established. Areas in which vegetation diversity declines due to establishment of new plant species often have lower insect diversity (Bezemer et al. 2014), which is consistent with our findings. Because *A. germinans* creates monocultures, we anticipate significant changes of coastal insect communities and a lowering of insect diversity and biomass, which may have large effects on ecosystem processes.

Insect community structure in GOM wetlands containing *A. germinans* was different from those reported from tropical mangrove forests, which tend to be dominated by Lepidopteran species influencing pollination (Landry 2013) and herbivory (Menezes and Peixoto 2009), with Dipterans making minimal contributions (Simberloff and Wilson 1969; Murphy 1990; Veenakumari et al. 1997; Burrows 2003). In our study, Lepidopterans only constituted about 5% of the groups collected, while Dipterans were the most abundant (31%). However, our Dipteran total is lower than totals reported in other salt marsh communities (Luckett and Gray 1966; Cameron 1972). Similarly, Murphy (1990), and Burrows (2003), only found herbivorous insects from four orders in mangrove forests, compared to seven orders found in this study, and the 10–12 found in studies looking at salt marshes (Cameron 1972; Wu et al. 2009).

There are some similarities between the fauna collected in mangrove forests in the GOM and those collected from tropical climates. For instance, when comparing feeding guilds, we found that sap-sucking insects were nearly three times less abundant in sites with *A. germinans*. These results are consistent with tropical mangrove forests which also show a lower abundance of sap-sucking insects (Veenakumari et al. 1997; Burrows 2003). Similarly, the areas with *A. germinans* in this study and the mangrove forests sampled by Burrows (2003) and Murphy (1990) found very few Coleopterans, which are normally abundant in coastal marshes (Cameron 1972). Mangrove forests can show a high degree of insect specialization with little to no overlap of herbivorous species in adjacent habitats (Burrows 2003). In contrast, we did not see separate families in sites with and without *A. germinans*,

possibly indicating that *A. germinans* in the GOM are in a transition between the communities found in tropical mangrove forest and those found in salt marshes. Our sites were within 15 km of one another and insects were likely able to travel between sites, which may also explain why we did not see distinct families associated with each wetland type.

Wetlands with rare *A. germinans* contained a larger diversity of plant species (e.g., *S. virginica*, *B. maritima*, *S. alterniflora*) than those dominated by *A. germinans* monocultures. It is not uncommon for encroaching plants to eliminate existing vegetation by competitive exclusion leading to decreased plant diversity (Wu et al. 2009; Harvey et al. 2010; Quan et al. 2016). For example, in China, the area occupied by *S. alterniflora* increased from 260 ha (Chung 1989) to 112,000 ha (An et al. 2007) in only 15 years with this expansion coming at the expense of existing vegetation (Wu et al. 2009). The diversity of lower trophic levels exerts significant controls on the abundances of the consumers that utilize them (Murdoch et al. 1972; Siemann et al. 1998; Knops et al. 1999). Encroaching plants can also alter habitat structure as they become established, influencing the existing arthropod communities (Gratton and Denno 2005). Our sites were exposed to similar abiotic factors (salinity, temperature, precipitation, etc.), and thus, the higher plant diversity, accompanied by the more labile plant materials associated with *S. depressa*, *B. maritima*, and the changes in the habitat structure are likely primary contributors to the higher insect abundances and biomass found in sites with rare *A. germinans*.

Changing plant compositions can alter food web structures in coastal wetlands (Gratton and Denno 2005, 2006; Levin et al. 2006; Harvey et al. 2010); however, patterns of community change have varied widely. In coastal marshes in southeastern Australia, native plants (*Juncus kraussii*) are being replaced by a congener (*J. acutus*). This change is associated with a lower diversity and abundance of predators (Harvey et al. 2010). In the northeastern United States, *P. australis* has displaced *S. alterniflora* in many areas. Gratton and Denno (2005) found feeding guilds in marshes containing abundant stands of *P. australis* to shift from a roughly 2:1:1 predator-to-herbivore-to-detritivore ratio to a 1:1 detritivore-to-predator ratio, with changes being driven by altered plant composition and physical structure. In our study, herbivores were the dominant guild in both the *S. alterniflora* and *A. germinans* habitat with the *A. germinans* dominated wetlands generally having fewer predators. The one exception to this is the inverted pattern found in fall mangrove sites. While it is possible that some environmental factor exists to explain this, for example, inverted biomass pyramids are common in aquatic systems (Wang et al. 2009), it is more likely that the dearth of fauna collected in fall mangrove sites has skewed the data. For example, only 18 arachnids were

collected within these wetland types; however, this small number accounted for nearly 23% of individuals, and just under 52% of the total biomass.

Although mangrove encroachment may alter community and ecosystem traits, mangroves provide benefits including protecting coastlines from storm surges and rising sea levels (Comeaux et al. 2012), which salt marshes can do as well. Bianchi et al. (2013) found that *A. germinans* may reduce some effects of climate change by sequestering carbon. If the current trend of mangrove expansion continues, the increased area occupied by *A. germinans* would increase the overall levels of carbon sequestration in coastal wetlands (Bianchi et al. 2013), providing an invaluable carbon sink which could help reduce the levels of CO₂ gas in the atmosphere. Based on this information, Bianchi et al. (2013) suggests implementing *A. germinans* planting programs to help increase the amount of carbon storage available in coastal wetlands. While the increased ability to sequester carbon is desirable, the possible consequences that may arise from a transition to mangrove forests from salt marsh habitat must be considered.

Avicennia germinans monocultures are replacing more diverse vegetation assemblages, and their presence has been associated with changes in insect community structure and changes in aquatic species composition (Diskin and Smee 2017; Smee et al. 2017). These changes may prove detrimental to coastal wetlands and the nearby ecosystems that rely on them. For instance, terrestrial arthropod species comprise a significant proportion of the diet of commercially important fish species in estuaries (Gray et al. 2002; Romanuk and Levings 2003) which further enforces the importance of wetland vegetation in estuarine food webs (Morley et al. 2012). This habitat loss is even more alarming when considered with other studies showing *A. germinans* presence adversely affects marsh nekton and benthic communities as well (Lunt et al. 2013; Diskin 2016; Smee et al. 2017).

This study provides insights into ongoing changes that are occurring in GOM estuarine wetland insect communities as *A. germinans* replaces native wetland plants. Future studies are needed to further elucidate the affect expanding mangrove forests could have on coastal wetland communities and food webs and the mechanisms by which these changes are occurring. Information gained would allow better action in the construction and implementation of future conservation policies and the management and restoration of coastal wetlands.

Acknowledgements Funding was provided by the USDA Forest Service Southern Research Station agreements 12-DG-11330101-096 and 13-CA-11330140-116 to D.L. Smee. The NSF-MSP ETEAMS Grant #1321319 provided funding for boat time and their interns, E. Urban in particular, assisted in the field. Members of the Marine Ecology Lab, and C. Trettin, J. Arnold, and C. Stringer from USFS provided

important assistance in the field. L. Patrick helped with manuscript formatting and proofreading. S. Bock was instrumental in writing and data analysis.

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