

Disease dynamics and potential mitigation among restored and wild staghorn coral, *Acropora cervicornis*

The threatened status (both ecologically and legally) of Caribbean staghorn coral, *Acropora cervicornis*, has prompted rapidly expanding efforts in culture and restocking, although tissue loss diseases continue to affect populations. In this study, disease surveillance and histopathological characterization were used to compare disease dynamics and conditions in both restored and extant wild populations. Disease had devastating effects on both wild and restored populations, but dynamics were highly variable and appeared to be site-specific with no significant differences in disease prevalence between wild versus restored sites. Disease affected up to 80% of colonies at one site following a tropical storm. A subset of 20 haphazardly selected colonies at each site observed over a single field season revealed widely varying disease incidence, although not in a consistent way between restored and wild sites, and a case fatality rate of 8%. Lastly, two field mitigation techniques, (1) excision of apparently healthy branch tips from a diseased colony, and (2) placement of a band of epoxy fully enclosing the diseased margin, gave equivocal results with no significant benefit detected for either treatment compared to controls. Tissue condition of associated samples was fair to very poor; unsuccessful mitigation treatment samples had severe degeneration of mesenterial filament cnidoglandular bands. Polyp mucocytes in all samples were infected with suspect rickettsia-like organisms; no bacterial aggregates were found. Overall, results do not support differing disease quality, quantity, dynamics, or health management strategies between restored and wild colonies of *A. cervicornis* in the Florida Keys.

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8 **INTRODUCTION**


9 Disease, in conjunction with co-occurring stressors such as storms and warming
10 temperatures, is the major driving factor placing the Caribbean staghorn coral, *Acropora*
11 *cervicornis*, at risk of extinction (reviewed in Aronson & Precht 2001, IUCN 2012).
12 Understanding the diagnostics and etiology of diseases affecting *A. cervicornis* populations
13 remains problematic, and effective management strategies to combat this ongoing threat to
14 species survival remain elusive. Despite more than a decade of focused research effort, there
15 remains a dearth of strict diagnostic characterization for field cases of disease in *A. cervicornis*.
16 Most authors simply apply the historical label of white-band disease (Gladfelter et al. 1977,
17 Peters 1984) or the more general label rapid tissue loss to what is likely a range of disease
18 conditions (Williams & Miller 2005, Raymundo et al. 2008). Corallivores, such as the snail
19 *Coralliophila abbreviata*, the polychaete *Hermodice carunculata*, and damselfishes or
20 butterflyfishes, further confound the ability to accurately assess disease by removing *A.*
21 *cervicornis* tissue and leaving feeding scars that may be difficult to distinguish from disease

22 (Bruckner 2001, Sutherland et al. 2004, Miller & Williams 2006). Further, Williams and Miller
23 (2005) found that *C. abbreviata* that had been feeding at tissue loss margins on disease-affected
24 colonies could transmit the condition to apparently healthy branches; thus predation may
25 exacerbate disease spread through a population.

26 *Acropora cervicornis*' status under the USA Endangered Species Act carries a legal
27 mandate to orchestrate its recovery (i.e., attain a sustainable status where ESA protections are no
28 longer needed to prevent extinction). This mandate, combined with a growing consensus that
29 decline has reached a point where natural resilience has likely been compromised, has led to
30 increasing efforts to culture and restock populations of *A. cervicornis* (reviewed in Young et al.
31 2012). This unprecedented movement toward proactive intervention and population engineering
32 in a coral reef foundation species is occurring within a historical context of mixed success in
33 previous case studies in the fields of fisheries and wildlife management (Hilborn & Eggers 2000,
34 Carlsson et al. 2008, Champagnon et al. 2012). The primary concern for such an endeavor is the
35 potential for unintended introductions of deleterious genetic or health consequences within the
36 imperiled population or its ecosystem (Cunningham 1996, Baums 2008). For this reason, the
37 genetic status of imperiled coral populations, including *A. cervicornis*, has received increasing
38 attention in recent years and strides have been made in addressing the potential genetic risks of
39 culturing and restoring *A. cervicornis* populations, such as outbreeding depression or genetic
40 bottlenecks in cultured stocks (Baums et al. 2010, Hemond & Vollmer 2010).

41 Addressing potential health risks of transplanting *Acropora cervicornis*, on the other
42 hand, is much more challenging. While explicit risk assessment and risk management
43 frameworks have been proposed and applied in wildlife translocation projects, effective
44 application requires at least qualitative knowledge of pathogens, vectors, and susceptibilities
45 operating in the given species (e.g., Lenihan et al. 1999, Sainsbury & Vaughan-Higgins 2012).
46 The lack of effective diagnostic tools and robust etiological characterization for coral disease in


47 general, and in *A. cervicornis* in particular (Sutherland et al. 2004, Rogers 2010), impairs
48 efficient health risk management. Until a better knowledge base is built for health management
49 of coral populations, general risk-averse best practices are currently conducted in *A. cervicornis*
50 culture and outplanting programs, including focus on field-based (rather than land-based) culture;
51 avoiding outplanting of colonies with visual signs of ill health (discoloration or tissue loss);
52 geographic restriction of source population, nursery site, and recipient populations; and focusing
53 outplanting at sites where there is evidence of prior occupation, but where no live wild colonies
54 currently exist (Johnson et al. 2011; L Gregg, Florida Wildlife Conservation Commission, pers.
55 comm.) 


56 The severe and ongoing impact of coral disease on coral populations begs the question of 
57 potential mitigation actions that could be applied in the context of local management (Bruckner
58 2002, Raymundo et al. 2008, Beeden et al. 2012). If effective, such targeted mitigation actions
59 would seem particularly relevant and useful as part of an integrated health-risk management
60 component in a population restocking program. Both nursery and field practitioners have
61 anecdotally reported simple interventions, such as separating apparently healthy tissues from
62 diseased colonies or applying a physical barrier (e.g., band of clay or epoxy) to the diseased
63 tissue margin to control tissue loss (Raymundo et al. 2008, Johnson et al. 2011), but no controlled
64 tests of such mitigation treatments have been published.

65 The objectives of the present study were to (1) characterize disease dynamics using
66 targeted disease surveillance in outplanted/transplanted versus wild populations of *A. cervicornis*
67 to provide a more robust scientific basis for judging the health risks associated with outplanting
68 and, (2) perform controlled tests of two simple mitigation treatments *in situ* to determine if they
69 significantly arrested tissue loss in affected colonies. For both objectives, and to improve our
70 understanding of the tissue loss diseases in this species, the histopathology of selected fragments
71 from unmanipulated and treated branches was evaluated using light microscopy.

72 MATERIALS AND METHODS


73 Study Sites

74 Disease prevalence surveys and mitigation treatments were conducted at restored and wild
75 *A. cervicornis* populations in the upper Florida Keys National Marine Sanctuary. Restored
76 populations were outplanted between 2007 and 2011 by the Coral Restoration Foundation (CRF)
77 or the National Marine Fisheries Service-Southeast Fisheries Science Center and initially
78 included a finite number (i.e., 18–50, some sites have proliferated considerably) of outplanted
79 (i.e., from field nursery culture) or transplanted (i.e., from nearby wild populations) colonies. 
80 These restored sites were deliberately established in areas devoid of native wild colonies and are
81 in shallow (3–8 m) fore-reef habitats, including Key Largo Dry Rocks, French, Molasses, Pickles,
82 and Conch Shallow reefs (Table 1; Supplementary Fig S1). An additional restored site
83 (Aquarius) was sampled in 2011 only and was located in the deeper fore-reef (14–16 m) of Conch
84 Reef. Few wild *A. cervicornis* patches are extant in the upper Florida Keys; three were identified
85 for the current study to provide comparison to the restored populations. These wild sites were all
86 located in low-relief patch reefs with partially consolidated rubble bottom at about 5 m depth and
87 included an unnamed patch reef off of Tavernier, FL (TavPatch sites A & B), and Little Conch
88 reef. Periodic surveys were also conducted at the CRF field nursery (origin of most restored
89 colonies).

90 Temperature data was collected at surveyed reefs during the survey seasons with HOBO
91 pendant data loggers (UA-001-64; Onset Corporation). Loggers were not re-located after
92 Tropical Storm Isaac in 2012 at TavPatch or Key Largo Dry Rocks  so no temperature data for
93 those two sites in 2012 is presented.

94 Disease Characterization


95 The primary disease reported to affect *Acropora cervicornis* is termed white-band disease
96 (WBD) (Gladfelter 1982, Peters et al. 1983, Peters 1984), and the tissue loss pattern resembling
97 that seen in the 1970s is now known as type I (WBD-I), because a type II was recognized in the
98 1990s, ~~WBD-II~~ (Peters 1984, Ritchie & Smith 1998, Aronson & Precht 2001, Vollmer & Kline
99 2008). ~~The~~ WBD-I disease was first reported from Tague Bay, St. Croix, US Virgin Islands, and
100 Gladfelter (1982) characterized this disease in *A. palmata* as “a sharp line of advance where the
101 distally located, brown zooxanthella-bearing coral tissue is cleanly and completely removed from
102 the skeleton, leaving a sharp white zone about 1 cm wide that grades proximally into algal
103 successional stages....” This is illustrated in Fig 1 A. Peters et al. (1983) found the same disease
104 signs present on *A. cervicornis* colonies of the deeper forereef at Tague Bay. WBD-I has now
105 been reported to occur throughout the Caribbean Sea (Aronson & Precht 2001, Raymundo et al.
106 2008), and is present in the Florida Keys (Fig 1 B). In WBD-II, a band of bleaching tissue 2–20
107 cm wide is present at the tissue loss margin, and its distribution had been limited to the Bahamas
108 (Ritchie & Smith 1998), although more recently seen in Puerto Rico (Gil-Agudelo et al. 2006)
109 and southeast Florida (EC Peters, unpubl).


110 The lesions resulting from tissue loss attributed to WBD on Caribbean acroporids have
111 varied in their patterns (smooth or ragged tissue margins) and rate (less than 1 mm d⁻¹ to more
112 than 14 mm d⁻¹, Gladfelter 1982), and descriptions of the lesions have not always been clear
113 (Rogers 2010). For example, “rapidly advancing white band of diseased tissue” (Vollmer & Kline
114 2008) is not appropriate because it is a band of white denuded skeleton, not white tissue, that
115 appears progressively (does not itself advance) from the base or middle of a branch toward the
116 branch tip as the necrotic tissue (confirmed by histological examination) peels off, sloughs, or
117 lyses and disappears from the skeleton (Gladfelter 1982, Peters et al. 1983). However, recent
118 observation of acute tissue loss in *A. cervicornis* in the Florida Keys indicates that disease rarely
119 present , a fairly uniform-in-width band of denuded skeleton beginning at the base of the

120 colony or more rarely in the middle of a branch, ~~as for~~ WBD-I (Fig 1C). Rather, initial lesions
121 often show irregular sloughing of tissue with rapid enlargement of lesions anywhere on the
122 surface of a branch, yielding multifocal swaths of bright white denuded skeleton referred to as
123 rapid tissue loss (RTL) (Fig 1D, Williams & Miller 2005). A similar but unnamed condition was
124 described by Bak and Criens in the early 1980's (1980). In the current field surveys, bright white
125 bare skeleton, either encircling a branch or in irregularly shaped patches, and adjacent to either a
126 straight tissue margin (distinct, smooth to undulating) or a jagged margin (distinct, serpiginous)
127 of sloughing tissue was classified as disease, WBD or RTL, respectively (Fig 1B-D). However,
128 sometimes tissue loss on a colony can appear as a combination of lesion types (Fig 1E). Rarely,
129 WBD-II was noted during this study (Fig 1F). Lesions where corallivorous snails (*Coralliophila*
130 *abbreviata*) were present as well as lesions confined to branch tips, but not past a fork, were
131 attributed to snail and fireworm (*Hermodice carunculata*) predation, respectively, rather than
132 disease (Fig 1G-H). The key features of the types of tissue loss in *A.cervicornis* are compared in
133 Table 2.

134 Surveillance

135 Disease surveillance was conducted from May to November ~~periods~~ in 2011 and 2012 to
136 target the seasonal time-frame when acroporid disease was expected to be most active (Willis et
137 al. 2004, Williams & Miller 2005, K. Nedimyer, pers. comm.). Surveys were conducted
138 approximately biweekly in 2011 (total nine surveys) and monthly in 2012 (total seven surveys),
139 each taking 2–3 days to complete. At the wild sites, a circular plot (8-m radius at Tav Patch A
140 and B, 10-m radius at Little Conch) was marked with a center rebar stake and used to delineate
141 the study population for which prevalence was determined (i.e., percent of colonies in the
142 population that displayed signs of disease). Different plot sizes were used at the two wild sites
143 incorporate a minimum of 25 colonies. At restored sites, the sample population consisted of the
144 outplanted and transplanted colonies. The number of colonies tallied for individual site

145 prevalence ranged from 23 to 163 according to the number of colonies available and the extent of
146 search during a given survey. During each survey, every colony was recorded as either affected
147 or unaffected with acute tissue loss disease (bright white skeleton with either a straight or jagged
148 tissue margin on basal portions of the colony or multifocal  Corallivores were sometimes
149 present and lesions attributable to these predators (either denuded branch tips characteristic of
150 fireworm feeding or usually basal lesions with snails present) were not counted as disease.
151 Prevalence was calculated for each site for every survey and averaged for each site-by-year
152 combination. A two-way, fixed-factor ANOVA, with factors being site-type (restored versus
153 wild) and year (2011 or 2012) and sites as replicates, was conducted to determine if overall
154 prevalence varied significantly between restored and wild sites or years. For reference, disease
155 prevalence observations were also made during six surveys in 2011 and one in 2012 at the nearby
156 field nursery (Coral Restoration Foundation) from which all the outplanted colonies in the study
157 had originated.

158 To characterize disease incidence and mortality, 20 randomly selected colonies were
159 tagged at each site in May 2012. At each survey, tagged colonies were photographed and a visual
160 estimate of percent of dead colony surface, attributed as either predation, disease, or undefined,
161 was recorded. After the fifth survey, disturbance from Tropical Storm Isaac damaged or removed
162 several tagged colonies at most sites, thereby yielding observations of fewer than 20 colonies at
163 the sixth survey. To determine disease incidence (rate of new disease cases) over a survey
164 interval, each colony observed with active disease which had been observed as unaffected at the
165 previous survey was counted as a new disease case. Incidence was expressed as a proportion of
166 observed tagged colonies displaying new cases of disease since the previous survey and was
167 standardized per week  t-test was used to determine if the proportion of tagged colonies that
168 remained unaffected during 2012 differed between restored and wild sites

169 We estimated partial mortality based on cumulative increase in rough visual estimates of
170 percent dead on each colony that was observed with disease. We analyzed cumulative partial
171 mortality for all cases which occurred through survey five, and then we included cases which
172 were observed at survey six, to include the acute mortality following Tropical Storm Isaac.
173 test was used to compare the proportion of affected wild vs. affected restored colonies showing
174 severe cumulative partial mortality (defined as greater than 80%). We also tallied the case
175 fatality rate as the percent of cases (colonies which displayed disease signs during the course of
176 the observation period) which displayed complete mortality.

177 **Mitigation Treatments**

178 Two disease mitigation treatments were implemented to test effectiveness in arresting
179 tissue loss (Fig 2). The first treatment used a band of two-part marine epoxy (All-Fix Epoxy)
180 applied around the branch to cover the disease margin of an affected colony, presumably
181 functioning as a physical barrier over the tissue-loss margin. The second treatment involved a
182 complete excision of live, apparently healthy, tips of branches distal to a disease margin using
183 handheld wire cutters. The excised fragment was then reattached to the reef substrate with epoxy
184 at a distance greater than 1 m from the parent colony. These treatments are referred to as epoxy
185 band (EB) and excision (EX) (Fig 2B-C), respectively. Lastly, a control treatment consisted of a
186 cable tie placed at or near a tissue loss margin on the same colony as a reference point to detect
187 continued tissue loss (Fig 1C or 1D). To prevent potential contamination, nitrile gloves were
188 used when manipulating colonies and were changed when moving between affected colonies. All
189 equipment that came into contact with diseased colonies was rinsed in a 10% bleach solution
190 following each dive.

191 The design and setup for this experiment (e.g., sample size, timing, and placement of
192 replicates) were constrained by the availability of affected colonies with apparently active

193 disease. Due to permitting constraints in 2011, no experimental mitigation treatments were
194 performed on wild *A. cervicornis* colonies. In 2012, this stricture was lifted and treatments were
195 conducted on both restored and wild colonies. Distribution of experimental replicates among
196 sites and years is given in Table 1. Effort was taken to block treatments within the same colony if
197 it contained three or four (to include a histology sample) affected branches. However, this was
198 often not possible and so treatments were allocated sequentially to affected colonies as they were
199 encountered.

200 Rates of tissue loss in the observed disease conditions were rapid so all experimental
201 replicates were scored as either (1) continued or (2) arrested tissue loss at an interval of
202 approximately one month after the treatment was implemented, and each treated colony was
203 photographed to document disease progression. In some cases, corallivores were subsequently
204 observed on a treated or control colony and these replicates were excluded from analysis.
205 Proportion of replicates with continued tissue loss was compared among the three treatments
206 using Chi-Squared test (for each year separately and for the years pooled).

207 **Histopathology**

208 To better characterize the observed disease conditions, tissue samples were collected in
209 2011 from a subset of apparently healthy colonies (n=21, including n=1-4 from each site
210 collected in June or late September 2011), diseased colonies observed in the vicinity of the
211 surveys (n=12), and diseased samples collected from the colonies in the mitigation experiment
212 (n=11), ~~collected throughout the sampling season~~. In addition, two diseased samples were
213 collected from wild site Little Conch in 2012 to compare with the apparently healthy samples
214 collected at that site in 2011. Samples were removed by cutting a 5–10 cm portion of a branch
215 (i.e., tissue and skeleton) using handheld wire cutters and placed in a labeled 50-ml plastic
216 centrifuge tube. After surfacing, the sample was immediately immersed in a formaldehyde-based

217 fixative solution (Z-Fix Concentrate, Anatech, Ltd., 1:4 dilution in seawater). Sample tubes were
218 capped, kept at ambient temperature in the shade, and shipped to the Histology Laboratory at
219 George Mason University for processing.

220 Each sample was photographed and the images compiled into trim sheets. Samples were
221 trimmed into approximately 2-cm long fragments using a Dremel tool and diamond-coated tile-
222 cutting blade. The location of each cut was marked on the sample image on the trim sheet and
223 subsample numbers were assigned and marked on the trim sheet. Subsamples having a tissue loss
224 margin were enrobed in 1.5% agarose to trap material that might be present on the denuded
225 skeletal surface or in corallite or gastrovascular canal crevices. Subsamples were decalcified
226 using 10% disodium ethylenediaminetetraacetic acid (EDTA) at pH 7, changing the solution
227 every 24 to 48 hours. When completely decalcified, the subsamples were rinsed in running tap
228 water for about 30 minutes, trimmed into 2–3 mm slices and placed in cassettes, processed
229 through ethanols, cleared, and infiltrated with molten Paraplast Plus®, then embedded in
230 Paraplast Xtra® (Peters et al. 2005). Sections (5- μ m thickness) were mounted on microscope
231 slides, stained with Harris’s hematoxylin and eosin and Giemsa (Noguchi 1926) or other special
232 staining procedures, and coverslipped with Permount™ mounting medium.


233 The sections were examined with an Olympus BX43 compound microscope and
234 photomicrographs obtained with an Olympus DP-72 camera. Semi-quantitative data (Jago 1996)
235 were collected from each subsample based on relative condition (0=Excellent, 1=Very Good,
236 2=Good, 3=Fair, 4=Poor, 5=Very Poor) and severity or intensity of tissue changes from normal
237 (0=Within Normal Limits, 1=Minimal, 2=Mild, 3=Moderate, 4=Marked, 5=Severe) (see Table S
238 1). Histoslides of *A. cervicornis* and *A. palmata* collected from the 1970s in the Florida Keys (the
239 earliest tissue samples located, before tissue loss was reported from this region) were used to
240 develop the “within normal limits” criteria for general coral tissue condition and zooxanthellae
241 condition/abundance scores, six specific cell or tissue parameters of polyp health, bacterial



242 aggregates (Peters et al. 1983), and suspect rickettsia-like organisms (RLOs) (Casas et al. 2004;
243 CS Friedman, pers. comm.). ~~Only presence~~ was noted for hypertrophied calicodermis foci,
244 necrotic cell spherules, suspect virus-like particles (VLPs) (PL Blackwelder, pers. comm.),
245 apicomplexans (Upton & Peters 1986), and suspect ciliate predators. The developmental stage of
246 gonads was noted, if present. Mean scores for each sample were obtained (one or multiple
247 sections were made, especially if enrobed samples had been trimmed into four ~ 2–3 mm slices
248 for embedding; some sections did not contain enough tissue for scoring) and checked for quality.
249 Suspect RLO abundances were visibly higher in Giemsa-stained sections since it demonstrates
250 *Rickettsia* well (Noguchi 1926), thus, estimates based on those sections were preferentially used.
251 Descriptive statistics were calculated for the scored parameters in each group of samples
252 (apparently healthy, disease characterization, and mitigation treatments). Frequency distributions
253 of the scores were examined. Comparisons were made for the scored parameters between all
254 apparently healthy and diseased samples, successful and unsuccessful mitigation treatments, and
255 WBD- and RTL-affected samples using Student's t-tests and Mann-Whitney U-tests.

256 **RESULTS**

257 **Disease Dynamics**

258 Disease prevalence was highly variable and largely site-specific with no consistent
259 patterns between restored versus wild sites (Fig 3 A-D). In 2011, wild sites showed consistently
260 low prevalence with means of 1.5 to 4.4% during the survey period and a peak of approximately
261 13% at TavPatch B in late June (Fig 3A). Meanwhile, four of six restored sites showed generally
262 high disease prevalence (i.e., survey period means of 9–17% and max of 26–41%; Fig 3C)
263 particularly from July through early October, while the remaining two restored sites showed
264 consistently low prevalence throughout 2011 (i.e., Key Largo Dry Rocks and Conch Shallow had

265 2011 survey period means of 0.7 and 3.5% prevalence with one peak of 13%, lower or similar to
266 the wild sites; Fig. 3C). Intermittent observations within the field nursery site throughout 2011
267 yielded consistently low prevalence of 0–1.7%. In contrast, during 2012, Key Largo Dry Rock 
268 and Conch Shallow showed among the highest prevalence patterns with survey period means of
269 20% and peaks of 60–70% (Fig 3E). Little Conch (wild) consistently had the highest site
270 prevalence throughout the 2012 survey period (20–57% range, mean 35%; Fig 3B).

271 The temperature records indicate little temperature variation between sites during both
272 years (Fig 3E-F), suggesting that site-specific disease increases or outbreaks were not triggered
273 by temperature. Additionally, the accumulated thermal stress was greater in 2011 than in 2012,
274 but the survey-period mean prevalence was higher in all the  wild sites and four of six restored
275 sites in 2012 . However, the passage of Tropical Storm Isaac (26 Aug 2012) corresponded to a
276 ubiquitous spike in prevalence ~~and the survey period maximum prevalence observed~~ across all
277 sites (restored and wild). A two-way ANOVA using site means for each year showed a significant
278 effect of year ($p=0.032$) but not of site-type ($p=0.786$) nor the interaction ($p=0.237$). However, if
279 the post-storm prevalence surveys are excluded in 2012, no factors are significant, suggesting that
280 higher overall disease prevalence in 2012 was attributable to the acute effect of the storm.

281 Temporal patterns of disease incidence in 2012 are shown in Table 3 and further
282 emphasize the site-specific nature of disease dynamics in this population. Individual sites show
283 widely varying patterns of incidence, from persistent low incidence followed by a spike in the
284 fifth interval, following Tropical Storm Isaac (e.g., Pickles, TavPatch-A, TavPatch-B), to a
285 moderate level in the first three intervals followed by declining incidence (Molasses), to sites
286 with persistently high incidence from interval two (French, Little Conch), to sites with both an
287 early and a late peak (intervals two and six; Key Largo Dry Rocks)(Table 3).

288 Among the initial tagged population of 160 colonies in 2012, a total of 89 cases were
289 identified with a case fatality rate of 7.9%. The proportion of colonies that remained unaffected

290 throughout the study (non-cases, Table 3) was not significantly different between restored and
291 wild sites (t-test, $p=0.686$). Prior to the storm (up to survey 5; only $n=53$ cases occurred up to
292 this point), 52% of both restored and wild cases showed no detectable increment of partial
293 mortality (Fig. 4) and frequencies of cumulative partial mortality were very similar. When the
294 storm interval is included, disease-affected restored colonies had a significantly greater likelihood
295 of having severe ($>80\%$) partial mortality than affected wild colonies (Fig. 4; z-test $p=0.005$).

296 **Mitigation Testing**

297 Approximately 60–70% of control replicates in each year showed continued tissue loss
298 after one month (Fig. 5). In other words, around one-third of the replicates we thought to be in an
299 active diseased state based on gross visual signs were, in fact, dormant during the following one-
300 month period of observation. The proportion of experimental replicates displaying tissue loss
301 about one month after the treatment application did not differ significantly among EB, EX, and
302 Control treatments for either year analyzed separately (2011: $\chi^2=0.134$, $p=0.935$; 2012: $\chi^2=1.502$,
303 $p=0.472$) nor for both years pooled ($\chi^2=0.953$, $p=0.621$). However, the power of these tests is
304 very low (0.059–0.173) so negative results should be treated with caution. Treatment success rate
305 appears to be slightly greater in 2012, specifically suggesting more likely benefit from EX
306 treatments than from EB treatments (Fig. 5).

307 **Histopathological Observations**

308 Summary statistics for the apparently healthy samples, diseased samples for
309 characterization, and diseased mitigation samples are presented in Table 4. The apparently
310 healthy samples were in very good to fair condition, had more zooxanthellae in gastrodermal
311 cells, numerous mucocytes that were about the same height as the ciliated columnar cells of the
312 epidermis (Fig 6A), and intact cnidoglandular bands of the mesenterial filaments (Fig. 6B). A

313 third of the samples had minimal gaps in the calicodermis, mesoglea, and epidermis covering
314 costal ridges. The calicodermis toward branch surfaces was squamous to columnar, relatively
315 thick, and contiguous over the mesoglea; calicoblasts often showed plasmallema extensions on
316 their apical surfaces (toward the skeleton) and pale pink to clear cytoplasm (Fig. 6C). Deeper
317 calicodermis was squamous and the cytoplasm contained fine eosinophilic granules. None of the
318 samples contained bacterial aggregates, but almost all had mild to marked numbers of suspect
319 RLOs in mucocytes on polyp oral discs and tentacles (Fig. 6D) and in cnidoglandular bands of
320 the mesenterial filaments (Fig. 6E). Coccidia oocysts were seen in a couple of samples. Early
321 oocytes were found in two samples, but no spermaries were observed.

322 Generally, characteristics of the diseased tissue samples collected from restored colonies
323 at a range of sites and throughout the 2011 season included moderate to severe attenuation of the
324 epithelia and mesoglea, numerous hypertrophied mucocytes or reduced number of mucocytes in
325 the epidermis (Fig 6F), reduced numbers of zooxanthellae (but not entirely missing), and cells of
326 the cnidoglandular bands showed varying degrees of atrophy, loss, necrosis or apoptosis, and
327 dissociation (Fig. 6G). Moderate to severe costal tissue loss was noted, beginning in the apical
328 polyp and increasing toward the tissue loss margin. The calicodermis varied in thickness and
329 condition, but deeper and closer to the tissue loss margin was thinner, had fewer cells, and
330 calicoblasts lysed or sloughed off the mesoglea (Fig 6H); sometimes foci of hypertrophied
331 columnar calicoblasts with apical fine acidophilic granules were present at lysing tissue margins.
332 None of these samples had bacterial aggregates, but all had suspect RLOs in mucocytes of the
333 oral disc and tentacle epidermis, cnidoglandular bands, and infected mucocytes were also present
334 in gastrodermis lining the gastrovascular canals and mesenteries (Fig. 6I-J). Suspect RLOs filling
335 epidermal mucocytes were large and pleomorphic (Fig. 6H), whereas those in gastrodermal
336 mucocytes were usually coccoid (Fig. 6I) and those in cnidoglandular band mucocytes could be
337 either morphology; size of the RLOs also varied. Tissue loss margins displayed lysing coral cells


338 with vacuolation and necrosis or apoptosis of cells remaining on the skeleton and sloughing of
339 epithelial cells from mesoglea. Some agarose-enrobed samples had free-swimming ciliates
340 containing zooxanthellae on the denuded skeleton in 24% of samples, but were very rarely in
341 contact with tissue; ciliates without zooxanthellae were present in fewer numbers on 10% of
342 samples, but farther away from tissue remnants. In addition, circumscribed masses of necrotic
343 cell debris and zooxanthellae, in various states of further degradation and lysing, were present in
344 33% of the diseased samples. About 1–2 mm in diameter, they appeared to form as calicoblasts
345 surrounding gastrovascular canals released from the skeleton and mesoglea surrounded
346 gastrodermal cells or mesenterial filaments or epidermis fragments, trapping the degenerating
347 epithelial cells within, but eventually becoming permeable to seawater and breaking apart. All of
348 the diseased samples obtained from colonies used in the mitigation treatments had similar
349 pathological changes (Table 4). Early to mid-stage developing oocytes were found in 10% and
350 5% of the samples, respectively, but no spermaries were observed.

351 Evaluation of the frequency distributions of the data to determine normality revealed that
352 most parameters had a bimodal distribution, divided between the apparently healthy and diseased
353 tissues (Table S 2), so the distributions were further examined within these categories. For
354 example, Epidermal Mucocytes Condition had no overlap in scores, with apparently healthy
355 samples showing mostly mild changes and diseased mostly severe changes. Parameters with
356 minimal overlap included General Condition 100x, Zooxanthellae Condition 100x, Dissociation
357 of Mesenterial Filaments, Costal Tissue Loss, and Calicodermis Condition. Parameters with
358 broader frequency distributions of similar scores for both diseased and apparently healthy
359 samples included Mesenterial Filament Mucocytes, Degeneration Cnidoglandular Bands, and
360 Epidermal and Filament RLOs.

361 Comparison of the apparently healthy samples with all diseased samples (Fig. 7A)
362 revealed that all parameter scores were significantly different, except for Epidermal and Filament

363 RLOs ($p=0.165$ and 0.767 , respectively, t-test, Table S 3). Epidermal RLOs were judged to be
364 moderate to marked in severity; Filament RLOs were mostly judged to be minimal to marked in
365 severity in both groups. For the samples in the mitigation treatments (Fig. 7B), histological
366 parameters were significantly different in unsuccessful treatments only for Mesenterial Filament
367 Mucocytes and Degeneration of Cnidoglandular Bands ($p=0.0097$ and 0.017 , respectively, Mann-
368 Whitney U-test, Table S 3). Number of mucocytes in the filaments was markedly fewer in
369 samples from colonies where mitigation was not successful, in addition the filament epithelium
370 had moderate to severe atrophy, loss of cnidocytes and acidophilic granular gland cells, and
371 necrosis or apoptosis of remaining cells. Samples categorized as WBD or RTL in their patterns of
372 tissue loss (Fig. 7C) only differed in Epidermal RLOs scores ($p=0.031$, Mann-Whitney U-test,
373 Table S 3).

374 DISCUSSION

375 Surveillance of multiple wild and restored populations of staghorn coral in the Florida
376 Keys during two years emphasize the ongoing toll that disease takes on this endangered species 
377 Devastating disease outbreaks appear intermittently in both wild and restored patches that have
378 appeared healthy for a number of years. For example, colonies at all three wild sites and restored
379 colonies at Key Largo Dry Rocks appeared healthy with minimal partial mortality (mostly
380 attributed to fireworm predation) throughout the 2011 surveillance, but two of these four sites
381 (one wild, one restored) were devastated by disease in 2012. All apparently healthy and diseased
382 samples collected in both years were infected with a suspect *Rickettsiales*-like bacterium (Casas
383 et al. 2004) (Table 4, Fig. 7A). Although Casas et al. (2004) dismissed this microorganism as a
384 potential pathogen of staghorn corals because it was present in apparently healthy and diseased
385 samples, as well as other coral species, our histopathological examinations revealed that it infects

386 polyp mucocytes and alters the coral's mucous secretions without causing gross disease signs,
387 potentially increasing the susceptibility of the coral to other environmental stressors and tissue
388 loss. There is no evidence that disease dynamics nor histological characterization are different
389 between wild and restored colonies within the study population, which suggests that different
390 disease risk management would not be warranted.

391 The high rates of disease prevalence documented in these populations are not unusual as
392 overall average disease prevalences of more than 25% have been reported for individual site
393 surveys in Panama, Belize, Cayman Islands, St. Thomas USVI, Antigua, and Curaçao for *A.*
394 *cervicornis* (Vollmer & Kline 2008, Fogarty 2012) and for *Acropora* spp. (Ruiz-Moreno et al.
395 2012). Somewhat lower, but still substantial, average levels of *Acropora* spp. disease prevalence
396 (8–12%) are reported in multi-year, Caribbean-wide, general coral condition surveys (Marks &
397 Lang 2007, Ruiz-Moreno et al. 2012). In comparison, disease prevalence in acroporid corals
398 across three sites in the Great Barrier Reef was also reported in the range of 9–13% (Willis et al.
399 2004), while more extensive surveys in three years across the entire Indo-Pacific region indicate
400 an acroporid disease prevalence of around 4% (Ruiz-Moreno et al. 2012).

401 The existence of disease-resistant genets within *A. cervicornis* has been reported at a
402 frequency of six percent in a studied population of 49 genets in Panama (Vollmer and Kline
403 2008). Four of the restored populations surveyed in this study are in fact genotypically
404 depauperate, containing the same three genets, while the other two restored populations were
405 genotypically more diverse (Table 1). Colonies at the three wild sites have not been genotyped,
406 but multiple genets and high genetic diversity have been previously documented in wild
407 populations of *A. cervicornis* in the Florida Keys (Baums et al. 2010; Hemond and Vollmer
408 2010). Thus, it is likely that multiple genets were present in each of these sites as well. The
409 detection of potentially disease-resistant genets is extremely problematic. Among the three wild
410 sites, we might have surmised possible disease-resistant genets within these presumably

411 genotypically-diverse patches given low disease prevalence in 2011. However, all of the tagged
412 colonies at wild site Little Conch were observed with disease at some point during 2012 (Table
413 3). An important goal of Caribbean *Acropora* population enhancement strategies is the nursery
414 culture of stress-resistant genotypes or phenotypes in order to propagate ~~hardier restored~~
415 populations (e.g., Bowden-Kerby & Carne 2012). The current results showing 1) extreme
416 variation in disease manifestation over sites and years, and 2) generally lower manifestation of
417 disease within the nursery environment than in nearby reef outplanted populations, (despite
418 similar RLO infection levels), reveal a challenge in accurately identifying these hardier
419 candidates.


420 Similarly, the site-specific nature of both disease prevalence and incidence patterns (i.e.,
421 patchy but not spatially autocorrelated) challenges the hope of identifying specific environmental
422 triggers for disease, at least on the site scale. While no severe warm thermal anomalies occurred
423 during the duration of this study, accumulated thermal stress was greater in 2011 than 2012—
424 corresponding to mild bleaching observed in some wild colonies during September–October 2011
425 (none in 2012)—but not greater disease impacts. Our temperature records do not indicate
426 substantial differences on the between-site scale that could account for spikes in disease among
427 our sites at different times (both within and between years, Fig 3E and F). Previous and repeated
428 reports of *A. cervicornis* disease in the Florida Keys have occurred in late spring to mid-summer
429 (April–July; Williams and Miller 2005; K. Nedimyer pers. comm.; M. Miller, pers. obs.), not
430 coinciding with the seasonal temperature peaks which occur in September–October. The only
431 coherent spike in disease prevalence and incidence that was discernible across all sites
432 corresponded to the passage of Tropical Storm Isaac (Fig. 3), corroborating the hypothesis that
433 storm disturbance may be an important coral disease trigger (Knowlton et al. 1981, Bruckner &
434 Bruckner 1997, Miller & Williams 2006, Brandt et al. 2013).

435 The only significant difference we were able to discern between restored and wild
436 colonies was in the degree of partial mortality during the storm interval, with restored colonies
437 having greater partial mortality than wild colonies (Fig. 4). One limitation of the current study is
438 in the spatial confounding of the restored and wild sites, with the former restricted to more
439 exposed, mostly shallow fore-reef habitats and the latter in somewhat more sheltered patch reef
440 habitats. It is likely that this habitat difference accounts for the apparent greater vulnerability of
441 restored colonies to storm-associated disease mortality rather than any inherent characteristic of
442 the colonies.

443 Our mitigation tests did not detect any significant benefit, in terms of preventing tissue
444 loss over a four-week period, from either excision or epoxy-band treatment. However, high
445 variability in response of both treatments, as well as the controls, yielded low power in the
446 statistical tests. There was a greater suggestion of benefit from treatments conducted in 2012
447 (Fig. 5). One difference between years was the inclusion of wild colonies in 2012 (only restored
448 colonies treated in 2011), but only a few replicates and only at Little Conch. Fisher's Exact Test
449 for control vs. pooled treatments and considering only wild colonies was still not significant (but
450 closer at $p=0.06$).

451 Several other observations may affect the interpretation of the somewhat inconclusive
452 mitigation test. First, there was no hint of harm accruing to either treatment (Fig 5). However,
453 we commonly observed in circumstances of high disease prevalence, a 'successful' (i.e., at one
454 month assessment) excision or other areas on a successfully epoxy-banded colony might resume
455 tissue loss at a later time, suggesting a re-activation of disease. On the other hand, if treatment
456 replicates that were implemented at times and sites with high prevalence (i.e., >15%) are
457 excluded, the remaining replicates indicate significantly lower frequency of tissue loss for
458 treatments (especially excisions) vs. controls (X^2 test; $p=0.014$; see Table S 4). Our results and
459 observations suggest that if mitigation interventions are attempted, branch-tip excisions are more

460 likely to be successful. Histologically, tip tissue may be in better condition than that at the tissue-
461 loss margin and resources are directed toward the tips rather than bases in this species (Highsmith
462 1982). Also, mitigation appears to be more successful in isolated cases rather than in areas with
463 more disease. Unfortunately, conditions with <15% disease prevalence ~~were surprisingly rare,~~
464 ~~occurring~~ in only 31 of our 56 individual site surveys in 2012.

465 The histopathological examinations revealed several other reasons why mitigation
466 treatment success can vary, despite the challenges in assigning a semi-quantitative score to
467 observations because specific changes occurring in the coral tissues formed a continuum. The
468 only significant differences in scores between the successful versus the unsuccessfully treated
469 branches were the greater loss of mesenterial filament mucocytes and degeneration of the
470 cnidoglandular bands of the filaments in samples from colonies that had unsuccessful treatments.
471 The filament epithelium lines the free edges of mesenteries in the gastrovascular cavity below the
472 actinopharynx in the polyp; the specialized acidophilic granular gland cells of this epithelium
473 release enzymes to break down food particles (Raz-Bahat et al. In Prep). The number and size of
474 gland cells and mucocytes in the cnidoglandular band increase, whereas ciliated cells decrease,
475 aborally in normal *A.cervicornis* tissue. Cell loss, necrosis, and lysing indicate that the polyp is
476 no longer able to process particulate food in the gastrovascular cavity. In addition, although
477 zooxanthellae condition appears to remain unaffected until the host tissue is sloughing off the
478 skeleton, their numbers are reduced as the host condition deteriorates. However, due to our
479 inability to detect changes in coral pigmentation until zooxanthellae numbers are reduced by
480 more than 50 percent (e.g., Jones 1997), the tissue grossly appears to be intact and “normal,”
481 when it may not be so microscopically. The ubiquitous presence of the suspect RLO infections
482 suggests most, if not all, the *A.cervicornis* population’s health is compromised  thus, without
483 microscopic examination, it is difficult, if not impossible, to identify the “best candidates” for
484 mitigation treatment.

485 Exactly what the impact of the suspect RLOs is on the *A. cervicornis* colonies is
486 conjecture at this point, but based on the behavior of similar obligate intracellular bacteria, their
487 replication within host cells requires substantial energy (Fryer & Lannan 1994) resulting in tissue
488 atrophy and necrosis (Friedman et al. 2000, Sun & Wu 2004). Nutritional stress may be a primary
489 reason why the zooxanthellae are gradually lost and calicoblasts lyse (Weis 2008, Schoepf et al.
490 2013). The coral cannot maintain its tissues with the loss of these host and algal cells that are
491 crucial to its survival. Infected mucocytes eventually die and are released from the epithelium and
492 the coral may not be able to replace them. Loss of mucocytes means the loss of the coral's
493 protection against sedimentation and microorganisms, as well as heterotrophic feeding (Brown &
494 Bythell 2005, Ritchie 2006). Investigation of the pathogenesis of RLO infection is continuing,
495 noting that other bacteria (*Vibrio harveyi*, *Serratia marcescens*, unspecified) have been implicated
496 in the acute loss of tissue from Caribbean acroporids (Patterson et al. 2002, Gil-Agudelo et al.
497 2006, Kline & Vollmer 2011). Transcriptome analysis shows gene expression alterations in
498 immunity, apoptosis, cell growth, and remodeling in WBD (Libro et al. 2013); and multiple
499 pathogens may be involved or be different in specific cases requiring histopathological
500 examinations (Work & Aeby 2011). Bacterial aggregates first proposed to be the pathogen (Peters
501 et al. 1983) were not present in any of these samples and ciliates do not seem to be a major factor
502 in tissue loss in our study. Histologically, no differences could be discerned between WBD- and
503 RTL-affected colonies, suggesting that differences in the patterns of tissue loss are due to the
504 intensity and duration of suspect RLO infections or the identity of other stressors that trigger the
505 loss. Samples collected from the same colonies in this study are also being processed for
506 molecular characterization of the microbial communities associated with them at the diseased
507 margin and in apparently healthy tissue from diseased or unaffected colonies to help explain the
508 pathogenesis of tissue loss.

509 Overall, our results confirm the devastating toll that disease continues to have on both
510 wild and restored populations of Caribbean staghorn coral and suggests that wild and restored
511 populations display similar disease conditions, dynamics, and impacts. These results emphasize
512 the continuing need to understand and effectively address disease impacts in this species, as well
513 as discover methods and run experiments to try and determine a way to minimize tissue loss of
514 diseased colonies. Unfortunately, the straightforward mitigation treatments tested in this study
515 provided equivocal benefit. Given this situation, population restoration might be viewed as a
516 necessary but stop-gap recovery measure, particularly in light of the suspect RLO infections of
517 mucocytes in nursery and wild colonies. Additional assessments of factors affecting the staghorn
518 corals and their tissue loss diseases are needed, including pathogen interactions between the
519 stocks (Friedman & Finley 2003) and host genotype susceptibility (Vollmer & Kline 2008).

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668

669 **Figure Legends:**

670 Figure 1: Illustration of disease and predation conditions categorized in this study. A) Loss of
671 necrotic tissue from skeleton of *A. palmata* during WBD outbreak, Tague Bay, St. Croix, 1980 B)
672 Typical disease-affected colony with multifocal lesions of denuded skeleton, C) WBD-I, D)
673 initial stages of RTL, (E) Colony manifesting signs of both WBD-I (base) and RTL (tips), F)
674 WBD-II signs, G) fireworm predation with two older preyed tips (partially colonized by algal
675 turfs) visible, and H) snail predation scar on basal portion of branch (removed snails indicated by
676 arrow) .

677 Figure 2: Illustration of the treatments used in the mitigation trials. A) Excision (EX) of healthy
678 looking tips snipped from a nearby disease colony and re-attached to the reef, B) Epoxy band
679 (EB) surrounding the diseased tissue margin. One month later (C) this ‘successful’ EB replicate
680 shows no additional tissue loss and initial regrowth over the epoxy. Control treatments are
681 illustrated in Fig 1C and 1D.

682 Figure 3: Disease prevalence in *Acropora cervicornis* colonies in Wild (A and B) and Restored (C
683 and D) populations over two survey periods (May–Nov 2011 and May–Nov 2012). Dotted lines
684 indicate close passage of Tropical Storm Isaac in Aug 2012. Panels E and F show the
685 temperature records from the same sites and time periods.

686 Figure 4: Frequencies ~~of severity~~ of cumulative partial mortality in tagged diseased colonies
687 during the 2012 survey period before (A and B, Surveys 1-5, n=32 and 21, respectively) and after
688 (C and D, Surveys 1-6, n=51 and 27 respectively) passage of Tropical Storm Isaac at Restored
689 and Wild sites. ~~More cases occurred after the post-storm disease spike.~~ The bin labeled zero
690 includes colonies that accumulated less partial mortality than could be resolved in coarse visual
691 estimates.

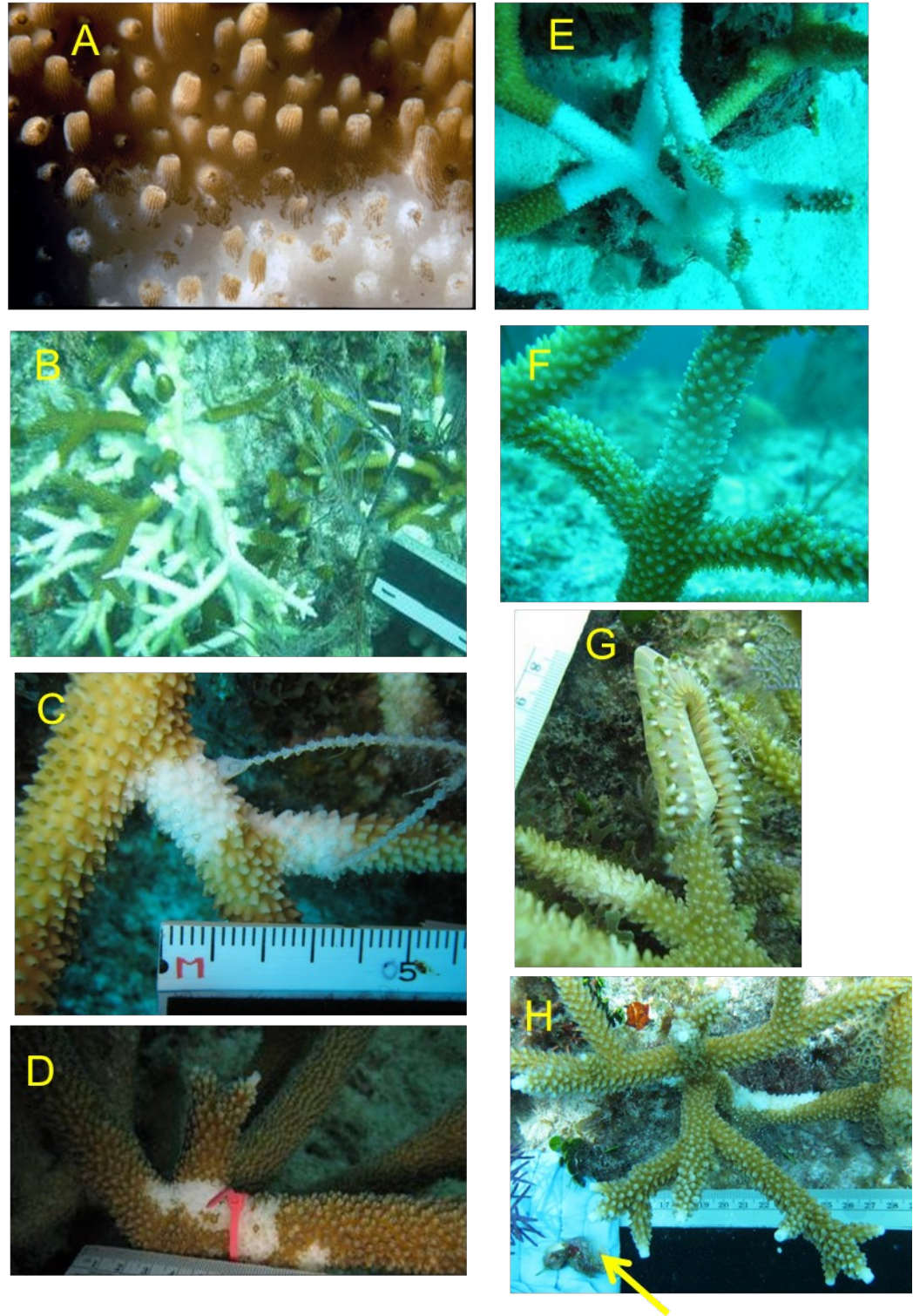
692 Figure 5: Results of experimental mitigation trials showing response in each year for Epoxy-
693 Band (EB), Excision (EX) and Control (cable tie placed around disease margin on a branch)
694 treatments as the percent of replicates showing continued tissue loss after one month. Number of
695 replicates ~~implemented given~~ above each bar. Chi-Squared Goodness of Fit tests indicate no

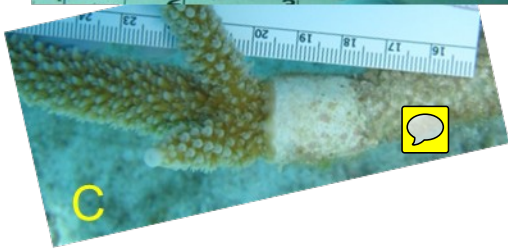
696 significant difference in the proportions of the three treatments showing continued tissue loss
697 when all replicates across years are pooled.

698 Figure 6: Histology observations. A) Coenenchyme epidermis from apparently healthy *Acropora*
699 *cervicornis* branch tip, columnar mucocytes of surface body wall (large arrow), suspect RLOs in
700 gastrodermal mucocytes of basal body wall (small arrows), Giemsa. B) Mesenteries showing
701 sections through cnidoglandular bands (large arrow), H&E. C) Apparently healthy staghorn
702 sample, epithelia lining gastrovascular canals with columnar calicoblasts having extensions of
703 plasmallema (large arrows), H&E. D) Section through tentacles (= T) and oral disc from
704 apparently healthy colony sample, mucocytes infected with suspect RLOs stain dark purple (large
705 arrow pointing to oral disc), Giemsa. E) Cnidoglandular bands from apparently healthy colony
706 sample, suspect RLOs in mucocytes (large arrows) and mucocytes in the epithelium (small
707 arrows). F) Coenenchyme epidermis from *A. cervicornis* showing signs of RTL, note atrophy of
708 epithelium and loss of mucocytes (large arrow), suspect RLOs in gastrodermal mucocytes of
709 basal body wall (small arrows), Giemsa. G) Sections through mesenteries from RTL-affected
710 sample with degeneration (necrosis, lysing) and dissociation of cells of the cnidoglandular bands,
711 note pink-staining acidophilic granular gland cells are rounding up and atrophied, ciliated cells
712 and mucocytes are reduced in number compared to Fig. 6B, H&E. H) RTL-affected sample
713 epithelia lining gastrovascular canals, severe atrophy of the calicodermis, loss of calicoblasts
714 from mesoglea (large arrows); adjacent gastrodermis is swollen, fragmented, and vacuolated
715 compared to cuboidal cells in upper left corner of image, H&E. I) Suspect RLOs infecting
716 gastrodermal cells (large arrows) lining the mesenteries (= MES) of an apparently healthy
717 sample, Giemsa. J) High magnification of infected epidermal mucocytes from apparently healthy
718 sample, showing pleomorphic suspect RLOs (large arrow) and mucocytes (small arrows, =
719 MUC), Giemsa.

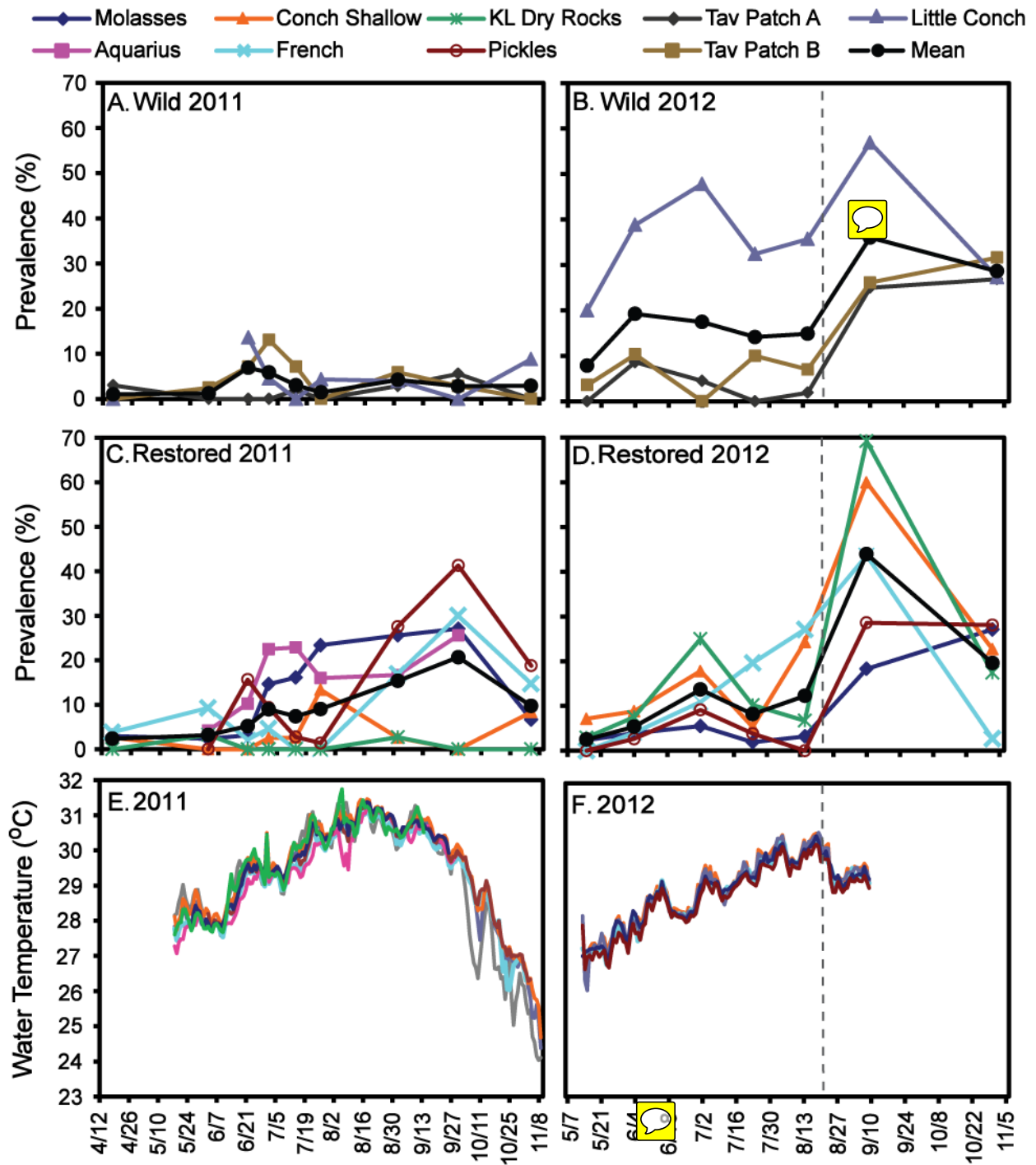
720 Figure 7: Histology parameter scores comparisons. A) Apparently healthy samples vs. diseased
721 samples. B) Successful vs unsuccessful mitigation treatment samples. C) Microscopic
722 characteristics of WBD vs. RTL samples.

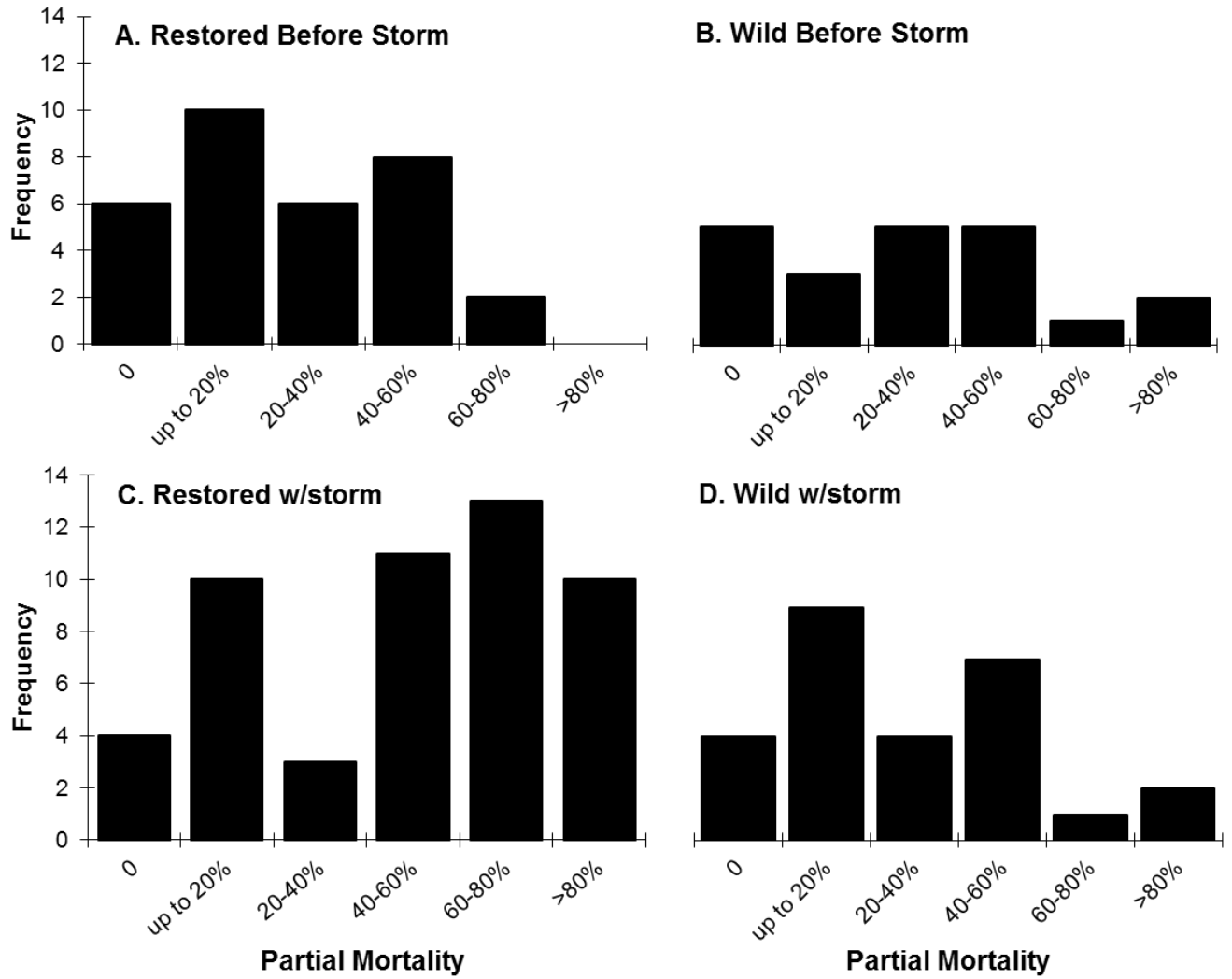
723 Figure 1:



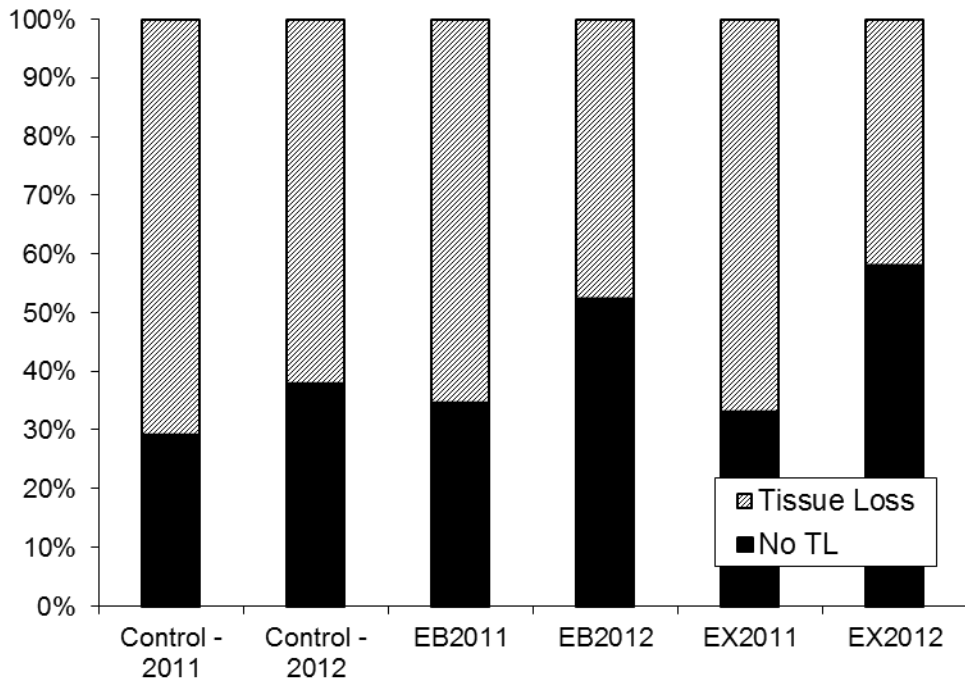


725 Figure 3:

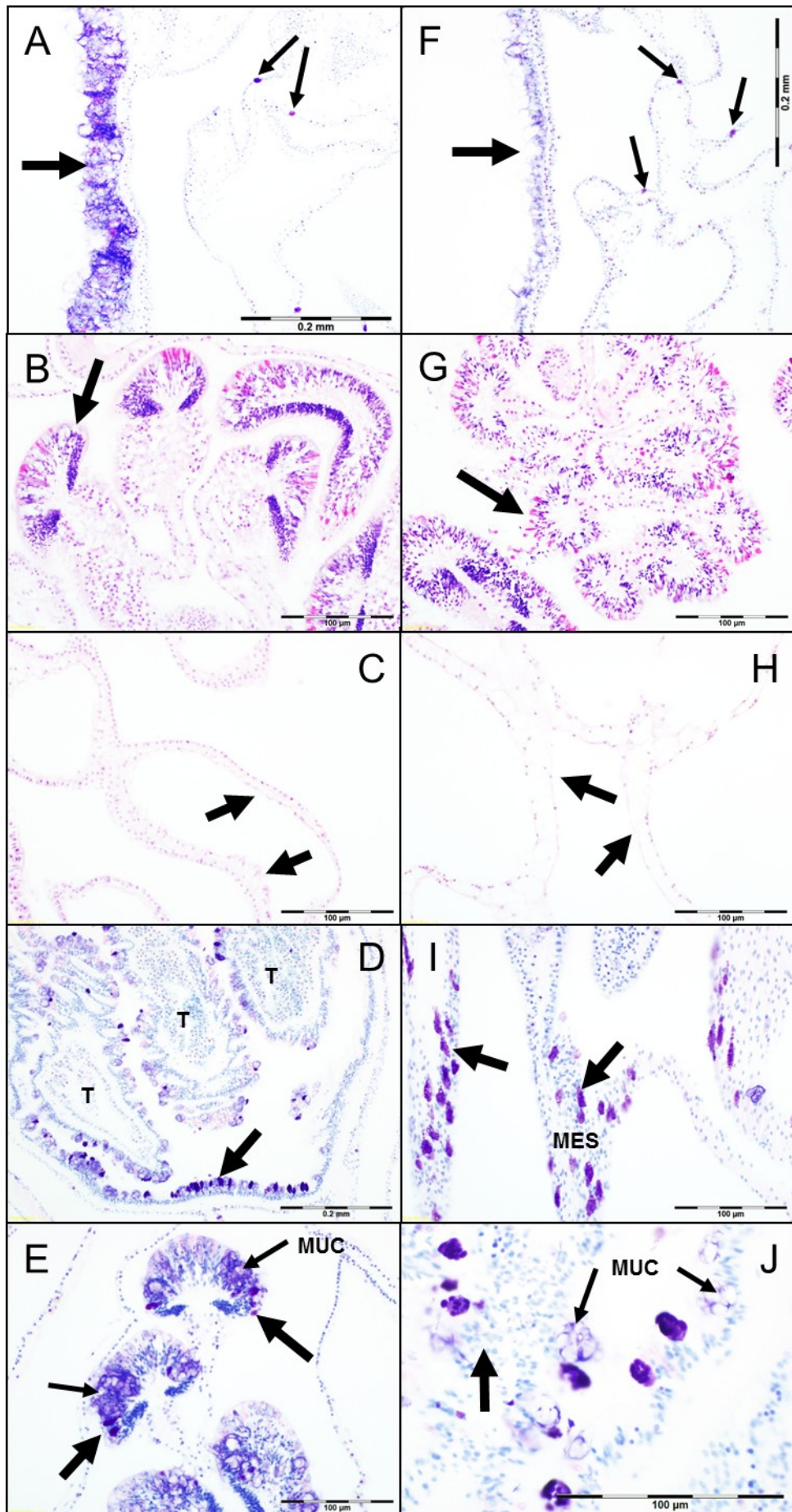




727 Figure 5:



728 Figure 6:



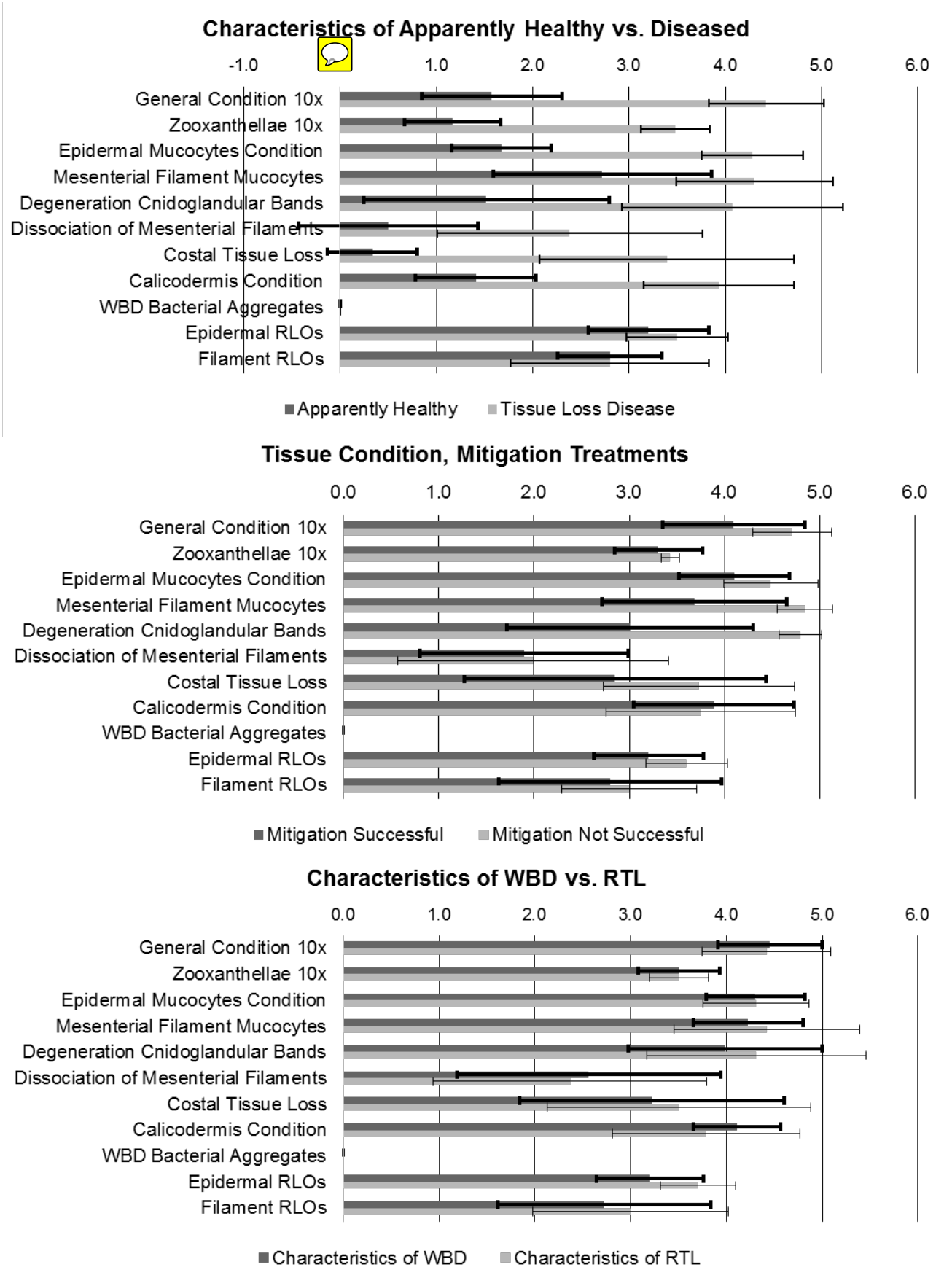


Table 1 (on next page)

Characteristics of study sites/populations in the upper Florida Keys

Number of genets indicates number of *Acropora cervicornis* multi-locus genotypes (based on seven microsatellite markers (Baums et al. 2009 , Baums et al. 2010)) within the surveyed populations at each site. Distribution of experimental replicates for the mitigation treatments among sites and years is summarized in the last two columns. UNK= Unknown, C=Control, EB = Epoxy Band, EX = Excision.

	Colony Origin	Site Type	# of Genets	Coordinates	Depth (m)	# 2011 replicates (C/EB/EX)	# 2012 replicates (C/EB/EX)
Molasses	Nursery	Restored	3	25° 00.60'N 80° 22.37'W	8-10	2/8/6	1/1/0
Aquarius	Transplant & Nursery	Restored	11	24° 57.20'N 80° 27.15'W	14	9/6/3	NA
Conch Shallow	Transplant	Restored	14	24° 57.08'N 80° 27.59'W	6	1/1/1	4/4/5
French	Nursery	Restored	3	25° 07.31'N 80° 17.85'W	10	5/3/4	6/5/1
KL Dry Rocks	Nursery	Restored	3	25° 07.45'N 80° 17.84'W	6	NA	3/3/4
Pickles	Nursery	Restored	3	24° 59.30'N 80° 24.74'W	8-10	0/1/1	NA
Tav Patch A	Wild	Wild	UNK*	24° 59.23'N 80° 27.17'W	6	NA	NA
Tav Patch B	Wild	Wild	UNK	24° 59.24'N 80° 27.16'W	6	NA	NA
Little Conch	Wild	Wild	UNK	24° 56.78'N 80° 28.21'W	6	NA	10/10/2
CRF Nursery		Nursery	>20	24° 59'N 80° 26'W	11	NA	NA

*Previous haphazard genotype sampling at this site yielded 6 unique genets in 20 sampled colonies (Miller & Baums, unpubl)

Table 2(on next page)

Disease Descriptions

Comparison of field manifestations of lesions seen in *A. cervicornis* and morphologic diagnoses. See Work and Aeby (2006) and Galloway et al. (2007) for definitions of terms.

Field Name	Tissue Loss Type	Location of Lesion on Colony*	Lesion Margin Appearance	Lesion Shape And Size	Lesion Number and Color	Lesion Progression	Morphologic Diagnosis
White-band disease type I (WBD-I) ¹	Acute to subacute	Base or middle of branch, encircling branch	Distinct areas of tissue loss, smooth to serpiginous margin, tissue tan to brown (due to symbiotic algae pigmentation)	Band of intact bare skeleton, well-differentiated from more distal skeleton	Focal to multifocal to diffuse, white (denuded skeleton), normally pigmented tissue margin	White band typically 2–10 cm wide; rate of tissue loss usually a few mm per day but can vary or stop; at branch bifurcation tissue loss continues on both branches at about the same rate; freshly denuded skeleton grades into green to brown algal growth on the skeleton, first visible after 5–7 days and becoming increasingly dense with time	Severe, basal to mid-branch band, diffuse, acute tissue loss, polyp, coenenchyme
White-band disease type II (WBD-II) ²	Acute to subacute	Tip or base of branch, encircling branch	Distinct areas of tissue loss, smooth margin, 2–20 cm wide band of bleaching tissue (loss of brown algal pigmentation) between tissue loss margin and normally pigmented tissue	Band of intact bare skeleton, well-differentiated from more distal skeleton, developing green to brown algal growth	Focal to multifocal, white (denuded skeleton), bleaching tissue margin	White band typically 2–10 cm wide; rate of tissue loss usually a few mm per day; bleaching margin tissue disappears, normally pigmented tissue starts bleaching; however, bleaching margin tissue may also disappear and then the normally pigmented tissue disappears, as in WBD-I; freshly denuded skeleton grades into green to brown algal growth on the	Severe, basal, band, diffuse, acute tissue loss, bleaching margin, polyp, coenenchyme

						skeleton, first visible after 5–7 days and becoming increasingly dense with time	
Rapid Tissue Loss (RTL) ³	Acute	Basal, medial, or colony-wide, partially to completely encircling branch	Distinct areas of tissue loss, undulating to serpiginous margin, tan to brown tissue, sloughing	Irregularly shaped areas of intact bare skeleton	Focal to multifocal and coalescing to diffuse, white (denuded skeleton)	Intact bare skeleton appears quickly along branches, new lesions may coalesce; rate of tissue loss usually cm per day; denuded skeleton develops green to brown algal growth that becomes uniformly visible after 5–7 days covering entire denuded area	Severe, basal to complete, band or irregular, diffuse, acute to subacute tissue loss, polyp, coenenchyme
Fireworm (<i>H. carunculata</i>) predation feeding scars ¹	Acute	Apical 1 to 5 cm of branch, but not extending beyond a branch bifurcation	Distinct areas of tissue loss encircling apex of branch, smooth to serpiginous margins, tissue tan to brown	Intact bare skeleton, tip of branch, developing green to brown algal growth	Focal to diffuse, white (denuded skeleton)	None, denuded skeleton develops uniform green to brown algal growth	Severe, focal, branch tip, acute tissue loss, polyp, coenenchyme
Snail (<i>C. abbreviata</i>) predation feeding scars ¹	Acute	Colony base, skeletal-tissue margin inward and vertically	Distinct areas of tissue loss, smooth to serpiginous rounded or scalloped margins, tissue tan to brown	Intact bare skeleton, usually adjacent to one or more <i>Coralliophila abbreviata</i>	Focal or multifocal, white (denuded skeleton)	None, denuded skeleton gradually colonized by green to brown algal growth	Severe, diffuse, basal, acute tissue loss, polyp, coenenchyme

*First lesion on all of these may be a single small focus of acute tissue loss, either at the base or in the middle of a branch, lesion enlargement pattern then varies.

¹Illustrated in Williams et al. (2006) but only for *A.palmata*

²Described in Ritchie and Smith (1998)

³Described in Williams and Miller (2005); described but not named in Bak and Criens (1981)

Table 3(on next page)

Incidence of Disease in 2012

Survey intervals (dates and duration in weeks), incidence, and proportion of colonies that remained unaffected by disease for the population of tagged colonies (n=20) at each site throughout the 2012 sampling period. Incidence is expressed as the proportion of new cases observed during each survey interval (i.e., diseased tagged colonies observed without disease in the previous survey) standardized per week. Shading is scaled with incidence value. Tropical Storm Isaac passed during Interval V.


Interval Dates (#weeks)	I 5/15-6/2 (2.71)	II 6/2-6/30 (4.00)	III 6/30-7.23 (3.29)	IV 7/23-8/15 (3.29)	V 8/15-9/10 (3.71)	 Unaffected	
Restored	<i>Conch Shallow</i>	0.018	0.000	0.000	0.064	0.126	0.400
	<i>Pickles</i>	0.000	0.025	0.000	0.000	0.094	0.400
	<i>Molasses</i>	0.037	0.025	0.000	0.016	0.000	0.800
	<i>French</i>	0.000	0.050	0.046	0.076	0.075	0.250
	<i>KL Dry Rocks</i>	0.018	0.088	0.000	0.015	0.184	0.150
Wild	<i>Little Conch</i>	0.037	0.063	0.046	0.076	0.099	0.000
	<i>Tav Patch A</i>	0.018	0.013	0.000	0.000	0.038	0.800
	<i>Tav Patch B</i>	0.018	0.000	0.016	0.015	0.058	0.700

Table 4(on next page)

Summary statistics for histopathological observations on all apparently healthy (n = 21), diseased (n = 11), and mitigation treatment samples (n = 11).

Parameter	Apparently Healthy			Characterization Diseased			Mitigation Treatments		
	Mean	St.Dev.	Range	Mean	St.Dev.	Range	Mean	St.Dev.	Range
<i>Assigned Scores</i>									
General Condition (100x)	1.6	0.7	1–3	4.5	0.5	3–5	4.4	0.7	3–5
Zooxanthellae (100x)	1.2	0.5	0–2	3.6	0.4	3–4	3.4	0.3	3–4
Epidermal Mucocytes Condition	1.7	0.5	1–2	4.3	0.5	3–5	4.3	0.6	3–5
Mesenterial Filament Mucocytes	2.7	1.1	1–5	4.4	0.7	3–5	4.2	0.9	2–5
Degeneration Cnidoglandular Bands	1.5	1.3	0–5	4.3	1.0	2–5	3.8	1.3	2–5
Dissociation Mesenterial Filaments	0.5	0.9	0–3	2.8	1.5	0–5	1.9	1.2	0.2–3.7
Costal Tissue Loss	0.3	0.5	0–1	3.5	1.3	1–5	3.2	1.4	0.9–4.8
Calicodermis Condition	1.4	0.6	1–3	4.0	0.7	2–5	3.8	0.9	2.1–4.9
Bacterial Aggregates	0.0	0.0	0–0	0.0	0.0	0–0	0.0	0.0	0–0
Epidermal RLOs	3.2	0.6	2–4	3.6	0.5	3–4	3.4	0.5	2.5–4
Filament RLOs	2.8	0.5	2–4	2.8	1.2	1–5	2.9	0.9	2–5
<i>Percent Affected (Presence/Absence)</i>									
Coccidia	10			14			10		
Calicodermis Repair	0			43			33		
Necrotic Cell Spherules	0			33			33		
Zooxanthellate Ciliates	0			24			24		
Non-zooxanthellate Ciliates	0			10			14		
Oocytes	10			10			5		
Spermaries	0			0			0		