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1 **Genet-specific spawning patterns in *Acropora palmata***

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14 Key Words: Florida Keys, Outcrossing, Synchrony, Fertilization potential, Temperature

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16 **Abstract**

17 The broadcast spawning elkhorn coral, *Acropora palmata*, requires outcrossing among
18 different genets for effective fertilization. Hence, a low density of genets in parts of its range
19 emphasizes the need for precise synchrony among neighboring genets as sperm concentration
20 dilutes rapidly in open-ocean conditions. We documented the genet-specific nightly occurrence
21 of spawning of *Acropora palmata* over eight years in a depauperate population in the Florida
22 Keys to better understand this potential reproductive hurdle. The observed population failed to
23 spawn within the predicted monthly window (nights 2–6 after the full moon in August) in three
24 of the eight years of observation; negligible spawning was observed in a fourth year. Moreover,
25 genet-specific patterns are evident in that 1) certain genets have significantly greater odds of
26 spawning overall and 2) certain genets predictably spawn on the earlier and others on the later
27 lunar nights within the predicted window. Given the already low genet density in this population,
28 this pattern implies a substantial degree of wasted reproductive effort and supports the
29 hypothesis that compensatory factors are impairing recovery in this species.

31 **Introduction**

32 Many broadcast spawning coral species, including *Acropora* spp., are obligate
33 outcrossers, requiring mixing of gametes from different genets for effective fertilization (Fukami
34 et al. 2003; Baums et al. 2005). Precise synchrony of gamete release among genets is crucial to
35 ensure successful larval production because broadcast gametes are viable over a period of only a
36 few hours and sperm dilution rapidly degrades fertilization potential in ocean conditions (Oliver
37 and Babcock 1992; Levitan et al. 2004). This required synchrony is expected to be particularly
38 important in populations with low densities of individuals and genets. *Acropora palmata*, the

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4 39 threatened Caribbean elkhorn coral, is genotypically depauperate in some areas of its range and
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6 40 significant declines in genotypic richness have been documented in the Florida Keys over very
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9 41 short time scales (Williams et al. 2014). These characteristics confer a particular need for
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11 42 spawning synchrony to allow for successful larval production in this population.

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14 43 *Acropora* spp. have less tightly constrained spawning nights than other genera of
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16 44 broadcasting corals. For example, in the Solitary Islands, Australia, Wilson and Harrison (2003)
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18 45 document a much wider range of spawning nights among individual colonies (including across
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20 46 months) for 12 *Acropora* spp. than among four mounding species. Among reports for Caribbean
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22 47 broadcasters, Fogarty et al. (2012) compiled spawning observations indicating that *Acropora*
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24 48 *palmata* spawns anywhere from night 1 to night 8 after the full moon (AFM), whereas Levitan et
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26 49 al. (2004; 2011) indicate that *Orbicella* spp. in both Panama and in the Bahamas show significant
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28 50 spawning on only two nights, within which individual corals did not show a preference for
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30 51 spawning on one night or the other.

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36 52 Currently, there are no published genet-specific records of spawning for Caribbean
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38 53 *Acropora* spp. This study documented the lunar nights of spawning by individual genets of the
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40 54 Caribbean elkhorn coral, *Acropora palmata*, in the Florida Keys over eight years to evaluate one
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42 55 aspect of successful larval production in this imperiled population.
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48 49 57 **Methods**

50
51 58 Spawning observations were made in the field during the expected *A. palmata* spawning
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53 59 window (nights 2–6 AFM, (Fogarty et al. 2012)) between 2130 and 2300 hrs (Eastern Daylight
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55 60 Time, representing ~1.5 to 3 h after sunset) during Augusts of 2007 to 2014. These observations
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57 61 were conducted at four sites in the upper Florida Keys National Marine Sanctuary, spanning a
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4 62 distance of approximately 18 km. Genotypes of all individual *A. palmata* colonies within study
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6 63 plots at three of the sites were previously determined and mapped from fixed landmark stakes
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9 64 (Williams et al. 2014). Observation effort was concentrated in the two reef sites with multiple
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12 65 genets present (Elbow and Sand Island reefs) though severe weather sometimes prevented
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14 66 planned observations. More opportunistic observations were collected at sites with a single,
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16 67 abundant genet (Horseshoe Reef, Baums et al. 2005; Molasses Plot 3, Williams et al. 2014) when
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19 68 additional boats and personnel were available. The area of observational coverage at each site
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21 69 was ~ 200–500 m², depending on the extent of *A. palmata* occurrence. Observations were made
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24 70 by divers patrolling a circuit among colonies within the observational area to examine polyps for
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26 71 ‘setting’ (the appearance of pink gamete bundles in polyp mouths) and subsequent release of
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28
29 72 gamete bundles. The *A. palmata* stands at Elbow, Sand Island, and Molasses reefs are typical of
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31 73 remnant fore-reef stands in the area ranging in depth from 3.5 to 4.5 m, whereas the monotypic
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33
34 74 stand at Horseshoe Reef is an atypically robust thicket occupying a nearshore patch reef (~ 3 m
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36 75 deep).

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38 76 At sites with only a single genet, divers simply observed and recorded substantial (i.e.,
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41 77 >50% of ramets spawning over a substantial portion of their live tissue) or minor (i.e., small
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43 78 patches of tissue releasing few bundles on only one or few ramets) spawning at that site. At
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46 79 diverse sites, genets were identified by arbitrary color names (e.g., Elbow Pink, Sand Island
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48 80 Blue). Divers used underwater maps and color-coded buoys to identify the genet of individual
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51 81 colonies when spawning was observed. Observers at these sites were all very familiar with the
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53 82 layout of colonies and genotypes at these permanent monitoring sites (Williams et al. 2008;
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55 83 Williams and Miller 2012). Observations were only recorded by night and genet; observations at
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58 84 the resolution of minute and ramets within genets were not consistently recorded as divers’

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4 85 attention was diverted to collecting bundles for other studies once spawning began. Bundle
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6 86 release from an individual colony in this species is protracted, generally spanning 15–20 min.
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9 87 A Kruskal–Wallis (nonparametric) ANOVA was used to test the hypothesis that the
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11 88 timing of spawning (i.e., median of any nights within the 2–6 AFM window when any degree of
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13 89 spawning was observed) differed significantly among genets. To further explore what factors
14
15 90 best predicted spawning success, an ordered logistic regression model otherwise known as a
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17 91 cumulative link mixed model was developed using the *ordinal* package in R (Christensen 2015).
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19 92 The degree of observed spawning was coded as an ordered response with no spawning < minor <
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21 93 substantial spawning. Year was included as a random effect and the two predictive variables,
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23 94 night after full moon (night AFM) and genet, were included individually and then together to test
24
25 95 which model best predicted spawning. Estimation of the predictive variables was via maximum
26
27 96 likelihood and the mixed models were fitted with the Laplace approximation (Agresti 2002).
28
29 97 Likelihood ratio tests were used to select the best model given the difference in the log likelihood
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31 98 score and the degrees of freedom of the models. P-values were calculated based on the chi-
32
33 99 squared distribution. Subsequently, this best fit model was used to calculate the maximum
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35 100 likelihood estimate and standard error of the ordered log odds of each individual genet spawning
36
37 101 compared to the genet with the least observed spawning (Elbow pink). The Z-score was used to
38
39 102 determine significance in these pairwise comparisons, with p-values less than 0.05 indicating
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41 103 genets with significantly greater odds of spawning on any given night. The exponent of the log
42
43 104 odds estimates for individual genets was calculated to obtain the proportional odds ratios for
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45 105 easier interpretation.
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53 106 Temperature loggers (HOBO Pendant) were maintained at Elbow and Molasses reefs
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56 107 over most of the eight years of the study, and at Sand Island since fall 2009, recording every 30
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4 108 min. These data were converted to daily averages and the mean of the three sites was graphed to
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7 109 visualize inter-annual variation in temperature regime between years.
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11 111 **Results**

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14 112 During three of the eight August spawning windows observed (2012, 2010, and 2007), no
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17 113 spawning by *A. palmata* was observed, and in a fourth year (2011), substantial spawning by only
18
19 114 two isolated genets was observed (Fig 1a; hereafter referred to as ‘poor’ spawning year).
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21 115 Observation during the September window only occurred in 2007, in which a single genet
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24 116 (Elbow gray) was observed to spawn. Observations were greatly curtailed in 2008 due to the
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27 117 occurrence of Tropical Storm Fay so spawning performance during this year is uncertain. Prolific
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29 118 spawning was observed in 2009, 2013 and 2014 with substantial spawning by many genets
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31 119 including co-occurring genets within sites (Fig 1a; hereafter referred to as ‘good’ spawning
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34 120 years) at a consistent hour (bundles released between ~ 2215 and 2240 hrs EDT). During these
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36 121 ‘good’ years, genets observed to spawn often did not spawn on the same night(s) AFM. However
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39 122 genets showed consistent patterns, spawning with highest probability either on early nights
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41 123 within the window (e.g., Horseshoe), on later nights within the window (e.g., Elbow green), or
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44 124 with little preference (e.g., Elbow orange; Fig 2). A one-way Kruskal–Wallis ANOVA
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46 125 confirmed significant variation among genets in the median night AFM of observed spawning (p
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48 126 = 0.03) though there was inadequate power to distinguish significant post-hoc pairwise
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51 127 differences among individuals.
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53 128 Based on single-factor ordered logistic regression models, night AFM and genet did not
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56 129 significantly differ in their ability to predict spawning. However, a model including both factors
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58 130 (Electronic Supplementary Material Table S1) showed significantly improved performance and
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131 displayed the lowest AIC score (Table 1). Additionally, three genets had significantly greater
132 odds of spawning on any given night relative to the least-spawning genet (Elbow pink) in
133 pairwise tests (Fig. 2). The genet with the greatest odds of spawning (46.33 greater than Elbow
134 pink) was the single genet at Horseshoe Reef. The genet at Molasses Reef and the orange genet
135 at Elbow Reef also displayed significantly greater relative odds of spawning (14.48 and 8.24,
136 respectively).

137 The temperature record, averaged among Elbow, Sand Island, and Molasses reefs during
138 the period of observation is given in Figure 1b. This temperature record shows no obvious
139 pattern of consistency among nor contrast between ‘good’ and ‘poor’ spawning years (Fig. 1).
140 For example, good years included ones with both relatively cold (2009) and relatively warm
141 prior winters while poor years also included cold (2010, 2011) and relatively warm (2007, 2012)
142 prior winters. Similarly, good years appear to have both relatively steep (2009) and less steep or
143 staged (2013) rates of warming during the first half of the year as do poor years (2010 vs 2012).

144
145 **Discussion**

146 Even during ‘good’ spawning years when multiple co-occurring genets spawned,
147 substantial spawning effort was observed on nights and at sites with presumably little potential
148 for fertilization success. Indeed, the model results suggest that the three individual genets with
149 the greatest spawning odds (Horseshoe, Elbow orange, and Molasses; significantly higher odds
150 than the ‘base genet’ of Elbow pink) each occupy a different site. For example, the monotypic
151 population at Horseshoe Reef is extensive (i.e., > 100 ramets; occupying >1500 m² of reef area),
152 comprised of colonies with little partial mortality which spawn profusely (MWM, pers. obs.).
153 However, it was observed to consistently spawn on earlier nights within the window (nights 2–4

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154 AFM; Fig. 2) than the more diverse sites (Elbow and Sand Island, predominantly nights 4–6
155 AFM; Fig. 2), or, in the case of 2011, when none of the diverse sites were observed to spawn at
156 all. In 2011, the Horseshoe genet did spawn the same night as the Molasses genet on night 4
157 AFM, but these sites are over 16 km apart, yielding extremely low expectation that these
158 gametes might mix at adequate concentrations within a few hours to effect fertilization.
159 Similarly, the orange genet at Elbow is also extensive (>30 ramets), displays little partial
160 mortality, and spawns profusely, but exerts a substantial portion of its spawning effort on nights
161 prior to most of its neighboring genets (e.g., 2009 and 2014, Fig 1a).

162 In the observed population, nights 5 and 6 AFM have the greatest genotypic diversity of
163 spawning colonies, implying greater fertilization potential (Levitan et al. 2004; Baums et al.
164 2013; Iwao et al. 2014). The degree of consistent genet-specific preference for spawning on early
165 (Horseshoe) versus later (Elbow white, Elbow green, and Elbow pink) nights within the observed
166 window seems in contrast to the report of no preference by individual *Orbicella franksi* for
167 spawning night within their smaller window (i.e., night 5 versus night 6 AFM; Levitan et al.
168 2011). Rather, genets of *O. franksi* have high hourly synchrony of spawning with proximity
169 among ramets as a secondary correlate of spawn time.

170 There are at least two possible explanations for the lack of observed spawning in certain
171 years: 1) corals may have spawned at a time outside our observations (a different month or lunar
172 nights outside the 2–6 AFM window); or 2) corals may not have spawned at all due to some
173 undescribed natural cycle or overarching stress that precluded physiological investment in
174 spawning (Szmant and Gassman 1990; Lirman 2000; Levitan et al. 2014). Summer 2011 is
175 recognized as a summer with warm temperature bleaching stress in the Florida Keys (Manzello
176 et al. 2015). Daily average temperatures exceeded 31 °C over ten days at the monitored

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4 177 spawning sites, with mild bleaching observed in *A. palmata* (DEW, pers. obs.). Thus, some
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7 178 degree of warm temperature stress in summer 2011 may correspond to lack of observed
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9 179 spawning in 2012, as shown in *O. faveolata* (Levitan et al. 2014). There was also a severe cold
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12 180 temperature stress event that especially affected nearshore Florida Keys reefs in early 2010
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14 181 (Lirman et al. 2011). However, this event did not appear to impact the *A. palmata* populations at
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17 182 the monitored spawning sites which are restricted to offshore habitats (Fig. 1b this study;
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19 183 Williams and Miller 2012).

21 184 It is perhaps more plausible that timing of spawning was simply displaced to months or
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24 185 nights beyond our observations during the ‘most likely’ window. Temperature is understood to
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26 186 be an important determinant of reproductive cycles in corals and thus changing temperature
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29 187 regimes with ocean warming have the clear potential to disrupt coral spawning phenology
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31 188 (discussed in Baird et al. 2009). Seasonal temperatures in the Florida Keys have already shown
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34 189 significant increase over historical baseline (Kuffner et al. 2015; Manzello 2015). However, the
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36 190 temperature record for three of the observed reefs over these eight years does not show obvious
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39 191 consistency among ‘good’ nor ‘bad’ spawning years (Fig. 1). There are some isolated reports of
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41 192 observed *A. palmata* spawning in months adjacent to our lack of observed spawning in August
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43 193 (e.g., July 2011 at Looe Key, D. Vaughn pers. comm.; Sept 2007; Fig. 1a) but did not appear to
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46 194 rise to the level of a full spawning event.

48 195 The rarity of spawning events in the observed population where multiple genets of *A.*
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51 196 *palmata* spawn in proximity on the same night indicate that larval production potential is even
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53 197 lower than the general characteristics of the population (e.g., colony and genotypic density;
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56 198 (Williams et al. 2014)) at these sites would suggest. It is anticipated that warming-induced
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58 199 phenological shifts (Baird et al. 2009), increased barriers to fertilization with ocean acidification
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4 200 (Albright et al. 2010; Albright and Mason 2013), and potentially other environmental stressors
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7 201 will only worsen processes of depensation in this population in the future. This prospect
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9 202 underscores the pertinence of proactive strategies to enhance both larval production and
10
11 203 successful recruitment of larvae as are now called for in the United States' *Acropora* spp.
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14 204 Recovery Plan (NMFS 2015) including potential transplants to increase density of synchronously
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16 205 spawning genets within reef patches.
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44 216 FKNMS-2012-101, FKNMS 2010-055, and FKNMS-2007-114).

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18 **281 Figure captions**

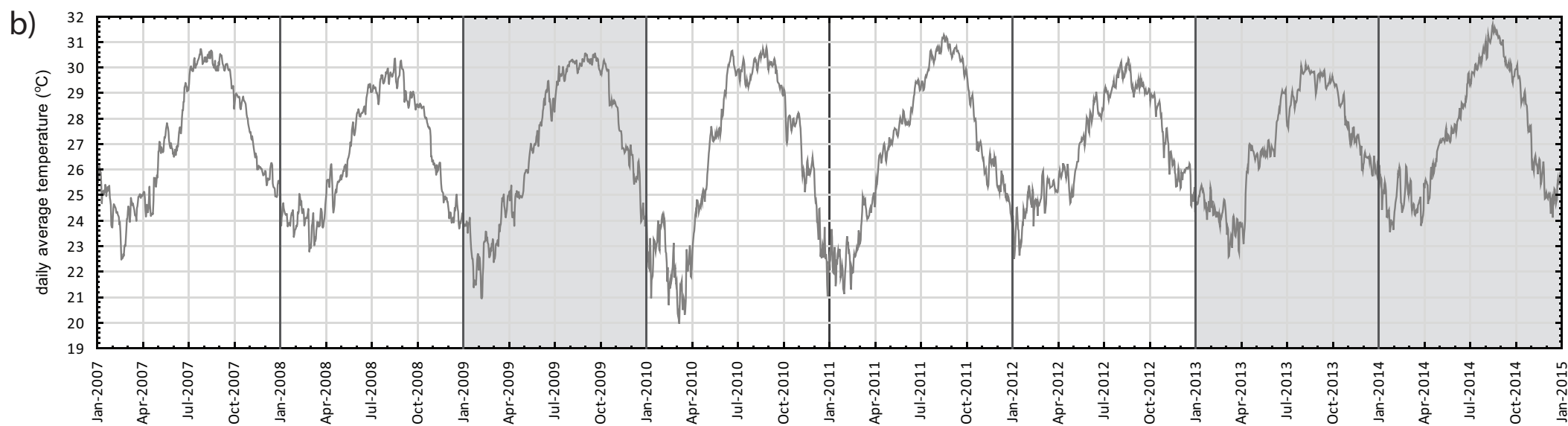
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22 **282 Fig. 1 a)** Spawning observations for twelve genets (depicted as rows) of *Acropora palmata*
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24 **283** across four reefs over eight years in the upper Florida Keys. *AFM* = after full moon. *Solid fill*
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26 **284** indicates significant spawning while *stippled fill* indicates minor spawning (see text for
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28 **285** characterization). *X* indicates no spawning observed. *Empty boxes* indicate that no observations
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30 **286** were conducted and *hatching* appears in a row to indicate the time when all colonies of that
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32 **287** genet within the observed reef area had died. *Asterisk* acknowledges that night 2 AFM for the
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34 **288** Sept 2007 lunar month was actually 31 August. **b)** Daily mean temperatures (based on in situ
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36 **289** HOBO Pendant loggers) averaged over three of the observed spawning sites (Elbow, Sand
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38 **290** Island, Molasses reefs) over the eight year spawning record. *Shading* highlights ‘good’ spawning
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40 **291** years, when multiple co-occurring genets were observed to spawn
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47 **292 Fig. 2** Frequency of spawning (number of nights observed) by individual *A. palmata* genets over
48
49 **293** the five-night expected spawning window pooled over eight years. Genets often spawn on more
50
51 **294** than one consecutive night. *AFM* = after full moon. P-values and proportional odds ratios (OR)
52
53 **295** depicted on each panel indicate significance of increased overall odds of each genet spawning on
54
55 **296** any given night (compared with the least-spawning Elbow pink genet) derived from the ordinal
56
57 **297** logistic regression model with night AFM and genet as predictive factors
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Figure 1

a)

		2007					2008					2009					2010					2011					2012					2013					2014																	
		Aug				31*	Sep				Aug					Aug					Aug					Aug					Aug					Aug																		
Night AFM		2	3	4	5	3	4	5	2	3	4	5	6	2	3	4	5	6	2	3	4	5	6	2	3	4	5	6	2	3	4	5	6	2	3	4	5	6	2	3	4	5	6	2	3	4	5	6						
Date		1	2	3	4	31*	1	2	18	19	20	21	22	7	8	9	10	11	26	27	28	29	30	15	16	17	18	19	3	4	5	6	7	22	23	24	25	26	12	13	14	15	16											
Elbow	Orange	X	X	X	X	X	X	X	Tropical Storm Fay											X				X	X	X	X	X		X	☐	X	X	X		X		X	X		X				X	X			X					
	Yellow	X	X	X	X	X	X	X											X	X	☐	☐		X	X	X	X	X	X	X	X	X	X	X		X	X	X	X		X				X					X	X	☐		X
	Gray	X	X	X	X	X													X	X	X	X	☐		X	X	X	X	X	X	X	X	X	X																				
	Blue	X	X	X	X	X	X	X											X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X		X		X	X		X				X	X				X	X			X
	Green	X	X	X	X	X	X	X											X	X	X	X	☐		X	X	X	X	X	X	X	X	X	X		X		X	X		X	☐			X	X	X		X					
	Pink	X	X	X	X	X	X	X											X	X	X	X	X		X	X	X	X	X	X	X	X	X	X		X		X	X		X	☐			X	X	X	☐	X					
	White	X	X	X	X	X	X	X											☐	X	X	X	☐		X	X	X	X	X	X	X	X	X	X		X		X	X		X				X	X	X		X					
	Sand Island	Orange	X	X	X	X	X	X											X	X	X	☐				X	X	X	X	X	X	X	X	X	X		X					X			X	X	X			X				
Blue		X	X	X	X	X	X	X	X	X	☐	☐			X	X	X	X	X	X	X	X	X	X		X					X			X	X	X			X															
Green		X	X	X	X	X	X	X	X	X	X	☐			X	X	X	X	X	X	X	X	X	X		X					X	X		X																				
Molasses										☐			☐		X	X	X	X	X	X	X			X	X		X	X	X						X					X														
Horseshoe																	X			X				X	X	X	X		X	X	X					☐	☐																	



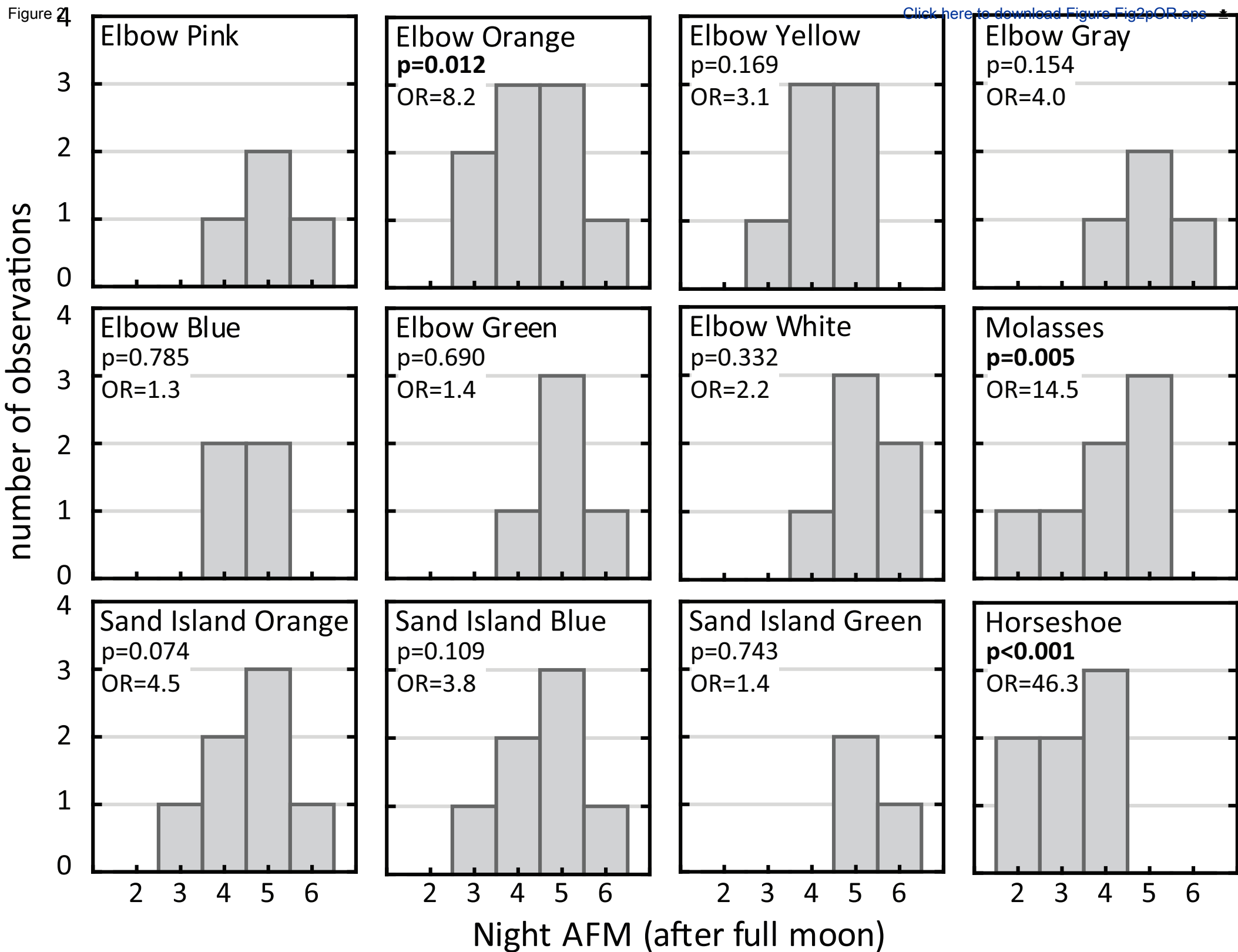


Table 1: Likelihood Ratio tests comparing ordered logistic regression models. All models include Year as a random variable. The likelihood ratio test compares the two models identified in the Test column and generates a p -value based on the difference in the $-2 \log$ likelihood statistic (Lik Ratio stat.) and difference in the degrees of freedom of the model residuals (Diff. in df) using the Chi-squared distribution. AIC indicates the Akaike information criterion for which lower values indicate better models.

Model	Predictive Variable	Residual df	AIC	$-2 \log$ Lik	Test	Diff. in df	Lik Ratio stat.	p -value (Chi-sq)
A	Night AFM	321	321.28	313.28				
B	Genet	311	339.08	311.08	A vs. B	10	2.20	0.99
C	Night AFM + Genet	310	313.10	283.10	B vs. C	1	27.98	1.22 e-07