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SPATIAL ECOLOGY AND CONSERVATION OF THE CENTRAL AMERICAN RIVER TURTLE (*DERMATEMYS MAWII*)

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SPATIAL ECOLOGY AND CONSERVATION OF THE CENTRAL AMERICAN RIVER TURTLE (DERMATEMYS MAWII)

A THESIS APPROVED FOR THE DEPARTMENT OF BIOLOGY

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ABSTRACT

Overexploitation of species is one of the primary causes of severe population declines. Conservation initiatives employ various strategies, such as protected land and captive breeding programs, to mitigate the effects of overexploitation. However, detailed knowledge of species' spatial ecology is a vital component of these successful conservation initiatives. We address the underlying need to elucidate spatial information for the Central American river turtle (*Dermatemys mawii*) — a wholly aquatic species that is suffering the effects of overharvesting to the point of endangerment— using two separate studies focusing on spatial ecology at broad and fine scales.

For the first study, we created a species distribution model to predict the extent of habitat suitability in Central America and quantify relevant factors that help define the ecological niche of *D. mawii* on a broader, regional scale. We examined overlap of highly suitable habitat and regions of increased hunting pressure represented by human settlements to better understand the optimal habitat with respect to environmental and anthropogenic factors. We expanded the potential range of *D. mawii* in Belize, Guatemala, and southern Mexico and found that areas of predicted suitable habitat were correlated with low elevation and moderate amounts of rain and precipitation seasonality. Although much of this area is fragmented, nearly one-half of this suitable habitat is outside zones of human settlement and may either represent previously unknown populations or be suitable for future attempts to bolster populations through translocations.

For the second study, we used radio telemetry to gain a better insight into the spatial requirements of *D. mawii* at a finer scale. We investigated juvenile home ranges at two separate sites on a single river in Belize. We calculated home range sizes using 95% and 100% minimum

xi

convex polygons and found the mean home range size (95% mcp) to be 3.53 ha. We determined that juveniles largely maintain a small home range, from which they occasionally travel, suggesting that protection areas may be sufficient to protect individuals.

Together, these two studies add to our understanding of the distribution and habitat requirements of this rare species. This information can serve to inform policy, conservation programs, and wildlife management practices.

1 Chapter 1: A Species Distribution Model for Conservation of the Central

2 American River Turtle (*Dermatemys mawii*)

3 ABSTRACT

The Critically Endangered Central American river turtle (Dermatemys mawii) is endemic to 4 Belize, Guatemala, and the Yucatan peninsula region of Mexico. Although this turtle is culturally 5 important, the exact range of this species is unknown. Defining the exact range is hampered by 6 7 the rarity and fully aquatic nature of this species. Here, we used species distribution modelling in 8 MaxEnt to extrapolate habitat suitability on a regional scale and estimate the importance of 9 several environmental predictors on the species' range. Our results show a limited area of high 10 habitat suitability in this region with approximately 50% of that habitat falling within areas of high-risk human impact, including hunting. Improved range and habitat suitability information 11 from this species distribution model is crucial to targeted conservation programs to protect 12 critical habitat areas and implement policy improvements. 13

14 INTRODUCTION

The Central American river turtle (Dermatemys mawii), the sole extant member of its family, is 15 found in lowland rivers and lagoons in the Yucatan peninsula region of Central America (Fig. 1) 16 (Iverson and Mittermeier, 1980; Vogt et al., 2011; Legler et al., 2013; Briggs-Gonzalez et al., 17 2019). Dermatemys mawii has historically been hunted for sustenance and remains culturally 18 important and highly sought after in this region (Moll 1986). Within the last 50 years, mass 19 20 harvesting practices and commercialization have caused extirpations and fragmented populations within its range (del Toro et al., 1979; Iverson and Mittermeier, 1980) leading to the IUCN 21 22 classification of Critically Endangered (Vogt et al., 2006). Beyond extirpation, direct effects

have not been documented in this species. Studies of other turtles and aquatic vertebrates show
that overexploitation by removing too many individuals from breeding populations cause further
population declines by decreasing clutch sizes (Daza and Páez, 2007; Eisemberg et al., 2017),
reducing recruitment (Walsh et al., 2006), and restricting genetic variability (Ratnerand Lande,
2001; Walsh et al., 2006; Allendorf et al., 2008). *Dermatemys mawii* populations are particularly
vulnerable to overexploitation by having long generation times characterized by low offspring
survivorship and long maturation times.

As D. mawii populations decline, conservation organizations are becoming increasingly 30 involved in programs to protect this endangered species by mitigating anthropogenic threats 31 32 while individuals grow and reproduce. Unfortunately, for many freshwater turtle species, merely allocating protected areas are inadequate to preserve endangered species populations (Norris et 33 al., 2019; Fagundes et al., 2016). However, populations in protected habitats can succeed when 34 35 combined with community involvement (Freitas et al., 2019), informed legislation (Karunarathna 36 et al., 2017), and captive breeding programs (Norris et al., 2019). Many such conservation strategies are dependent on identifying where a species is likely to occur. 37

To counter the serious threat D. mawii is facing and guide future field research, a more 38 thorough understanding of this species' distribution and the environmental variables that shape 39 its distribution is necessary (Sinclair et al., 2010). The limited knowledge of D. mawii's range 40 41 and ecological niche are based on locality data collected during diet studies (Legler and Vogt, 2013), population genetics studies (González-Porter et al., 2011, 2013), and habitat preference 42 observations (Legler and Vogt, 2013), with few surveys focused on understanding the full extent 43 44 of its distribution or fundamental niche (Vogt et al., 2011, Rainwater et al., 2012). A greater ecological understanding of this species, improved planning for reintroduction sites, and 45

46 programs for habitat protection and conservation can be achieved by integrating habitat

47 suitability information from species distribution models (Bombi et al., 2011; Guisan et al., 2013).

Species distribution models (SDMs) provide information on the fundamental niche of species 48 by combining specific environmental factors (Grinnell, 1917; James et al., 1984) and the 49 geographic distribution of species obserations (Hutchinson, 1959; James et al., 1984). 50 Unfortunately, both of of these data types are challenging to obtain for cryptic or endangered 51 52 species, and available data for *D. mawii* are scarce. Field surveys are often expensive and time intensive, whereas SDMs can be run using available locality data from prior studies. Machine 53 learning algorithms, like Maximum Entropy (MaxEnt), predict areas of habitat suitability using 54 55 presence-only occurrence points while remaining robust against small sample sizes (Anderson and Gonzalez, 2011; Guillera-Arroita et al., 2014), which often occurs with rare or endangered 56 species. In this study we i) extrapolate the extent of the suitable habitat of the Central American 57 river turtle using machine learning software, ii) determine the environmental variables that are 58 59 associated with the distribution, and iii) identify optimal regions for conservation efforts in relation to human impact. We predict that habitat suitability will be constrained by elevation and 60 rainfall. By using confirmed presence points and a combination of pertinent variables to train the 61 model, we can map areas of high habitat suitability and areas of human impact on D. mawii. 62 63 Using this map, we can predict the likelihood of occurrence of *Dermatemys mawii* in Central America as well as determine areas well-suited for conservation planning. 64

65 **METHODS**

66 **Presence points and predictor variables**

67	Locality coordinates were collected from previously published literature (Vogt et al., 2011;
68	Rainwater et al., 2012), unpublished surveys, and museum database searches using VertNet
69	(vertnet.org; Guralnick and Constable, 2010) for records containing Dermatemys mawii.
70	Presence data were compiled and georeferenced to ensure accurate coordinate data in ArcMap
71	(ESRI, ver 10.6.1). Sampling design for field surveys are assumed to be as randomly sampled as
72	possible from accessible rivers and lagoons, however, to avoid spatial autocorrelation and correct
73	for sampling bias in areas where multiple surveys occurred, presence points were systematically
74	subsampled in ArcMap within 1 km grids to reflect the resolution of the environmental variables
75	(Fourcade et al., 2014).
76	We obtained BIOCLIM environmental variable raster files from the CHELSA database (Karger
77	et al., 2017) averaged from 1979–2013 at a spatial resolution of 30 arc-seconds (1 km) in a
78	WGS84 projection (Table 1). We clipped these layers to the extent ($xmin = -100$, $xmax = -85$,
79	ymin = 12, ymax = 24) of our study region to include Guatemala and Belize, as well as portions
80	of Mexico, Honduras, and El Salvador. We also included topographical elevation data from
81	USGS GMTED2010, which is a suspected to be biologically relevant due to this species'
82	reliance on permenant water bodies. This layer was then resampled to the same cell size and
83	clipped to match the extent as the environmental variables in ArcMap (ESRI, ver 10.6.1).
84	To reduce collinearity and overfitting (Warren et al., 2014; Zeng et al., 2016) applied a principal
85	component analysis (PCA) to identify the factors that best explain the variance at the presence
86	points (Demšar et al., 2013). To further reduce collinearity, we used a Pearson's correlation test
87	to confirm that all environmental features were below a 0.7 cutoff value (Gogol-Prokurat, 2011).
88	Model parameters and assessment

We used the ENMeval package (v. 0.3.0; Muscarella et al., 2014) to set data-specific parameters 89 and run the model implementing the MaxEnt algorithm (Phillips et al., 2006) in R (R Core Team, 90 91 ver.1.3.1073). A minimum convex polygon hull was created to encircle the outer presence points, within which 10,000 background points (Muscarella et al., 2014) were randomly selected 92 for use in characterizing the environment. Model complexity was balanced by running models 93 94 across various regularization multipliers and feature classes: linear, linear quadratic, and linear quadratic product (Muscarella et al., 2014; Peterson et al., 2011). Regularization multipliers from 95 one to three at intervals of 0.5 were applied to the model to correct for overfitting (Merow et al., 96 97 2013). After models were completed, we determined the best fit model from the lowest Akaike information criterion (AICc) value, which corrects for small sample sizes (Warren and Seifert, 98 2011). We evaluated the importance of environmental variables using permutation importance 99 rather than percent contribution because permutation importance relies on the final model rather 100 than the paths used in a particular run (Songer et al., 2012). Models were run and averaged using 101 a 5-fold cross validation (Muscarella et al., 2014). The output habitat suitability map is a cloglog 102 raster, which rescales the habitat suitability scores from zero to one, where zero represents lowest 103 predicted suitability and one represents highest predicted suitability (Phillips et al., 2017). 104

105 Habitat calculation and human impact

We used the mean habitat suitability value from the MaxEnt output to determine the cutoff for
high habitat suitability (Liu et al., 2005; Cramer, 2003). We calculated the total amount of high
habitat suitability by multiplying the total number of cells by the cell size. To represent regions
of higher human hunting pressures, we included a human settlement polygon to the habitat
suitability map. We transformed the point layer GHS POP (Schiavina et al., 2019; Freire et al.,
2016) from the Mollweide projection to WGS84 to match the environmental layers and clipped it

to the output of areas of high habitat suitability. A 5 km radius buffer was then added to each

113 human settlement to represent the likely concentration of hunting using boats and freediving near

settlements (Peck et al., 2011) before merging all points into one polygon. We calculated the area

of human impact layer with the same procedure as the area of high habitat suitability and

determined the amount of overlap in suitable habitat with potentially high hunting pressure.

117 **RESULTS**

118 Species distribution model

119 We used a total of 68 locality points across Mexico, Guatemala, and Belize to train the model,

120 after data quality checks and subsampling highly clustered localities.

121 With regard to the PCA used to evaluate which of the factors best explain the variance at the

presence points, the first two principle components explainexplain 71.9% (PC1: 40.9% and PC2:

123 31.0%) of the variance. Precipitation of the wettest month, precipitation of the wettest quarter,

124 precipitation seasonality, and precipitation of the warmest quarter show the highest contributions

to PC1 and PC2 (Fig. 2).

126 The Pearson correlation test of the four variables selected from the PCA and the topographical

variable revealed high collinearity (0.995) between precipitation of the wettest month (Bio 13)

and precipitation of the wettest quarter (Bio 16), however all other variables were below the

cutoff value (0.7) (Table 2). We excluded Bio 16 due to the high collinearity and lower relativecontribution to PC1.

131 The habitat suitability model with the lowest deltaAICc score was determined to be the best fit

model for the data with an acceptable AUC value of 0.795 (Gogol-Prokurat, 2011; Hosmer et al.,

133 2013) using the linear quadratic product feature class at regularization multiplier 2.

134 Habitat suitability and human impact area

135 Elevation shows the highest permutation importance (57.2%) followed by precipitation

seasonality (27.3%), precipitation of wettest month (8.8%), and finally precipitation of the

137 warmest quarter (6.7%) (Table 3). At lower values, elevation, precipitation of wettest month, and

138 precipitation seasonality, habitat is predicted to be more suitable (Fig. 3). Conversely,

139 precipitation of the warmest quarter of the year predicts suitable habitat at higher values.

140 The overall habitat suitability values range from 3.67e-09 to 0.9518 across the study extent. The

141 mean habitat suitability value (0.213) was used for the threshold of high suitability. High

suitability habitats are located in lowlands of Mexico, Guatemala, Belize, and a small area in

143 Honduras covering approximately 281,722 km² (Fig. 4) The human impact area overlaps 51.6%

of this highly suitable area, leaving 136,000 km² of the environmentally suitable area with lower
human impact (Fig. 5).

146 **DISCUSSION**

Using MaxEnt for species distribution modelling, we set out to identify the extent of the suitable 147 habitat of *D. mawii*, ascertain environmental factors associated with its distribution, and identify 148 optimal regions for conservation efforts in relation to human impact. We determined that 149 climatic habitat suitability of *D. mawii* is over 281,000 km² in Central America, however, this is 150 not excluded solely to aquatic habitat. This identified suitable habitat notes highly suitable 151 152 habitat for translocations; and if the species is present in the contained aquatic habitats, would expand the distribution range known from existing literature (Vogt et al., 2011; Legler and Vogt, 153 2013). Specifically, the potential range provided from the SDM extends the current range to 154 include suitable habitat further north into the Yucatan Peninsula and into Honduras where D. 155 mawii has not been documented. The species distribution model also refines the potential 156

distribution by excluding areas of low suitability from the wider known distribution range in
Belize and Guatemala. The model indicated that low elevation, moderate amounts of rainfall in
the wettest month of the year, lower precipitation seasonality, and high precipitation in the
warmest three months of the year are important factors for habitat suitability, therefore providing
insight into the ecological niche of *D. mawii*.

High habitat suitability is most strongly associated with lowland habitats and sharply declines 162 163 with an increase in elevation. Rather than functioning as a physiological constraint, changes in elevation likely limits the availability of slow-moving, permanent streams and lagoons. Greater 164 amounts of rainfall are associated with high elevation in this region, therefore, areas with higher 165 166 rainfall show lower habitat suitability while conversely, low to moderate rainfall in both the wettest month of the year and warmest quarter of the year are correlated with higher habitat 167 suitability. Mild precipitation seasonality, or moderately consistent precipitation throughout the 168 year with a few months receiving more rainfall than the rest, is also important to their habitat 169 170 suitability given their fully aquatic nature (García-Anleu et al., 2011; Legler and Vogt, 2013; Briggs-Gonzalez et al., 2019) and apparent reliance on the inundation of floodplains for seasonal 171 dispersal (Briggs-Gonzalez et al., 2019; Legler and Vogt, 2013). 172

The SDM accurately predicts areas of high habitat suitability based on environmental variables, however, there are geographical constraints that are not considered in our model. The SDM shows high habitat suitability extending north into the Yucatan peninsula based on the chosen variables. However, the peninsular region is primarily karst, and is therefore unlikely to contain *D. mawii* due to the lack of surface water. Rainwater that does not evaporate is either absorbed into a layer of decomposed limestone or seeps into the aquifer below (Perry et al., 2003). Absence of major rivers or streams greater than 100 m in the northern Yucatan Peninsula

(Perry et al., 2003; Perez et al., 2011) likely prevents turtles from accessing regions of the 180 peninsula. Additionally, a lowland region in northern Honduras and a narrow stretch of land 181 182 along the Pacific coast of Mexico and Guatemala are indicated by the SDM as suitable habitat but are unlikely to contain *D. mawii*. The area of suitable habitat in Honduras is marginally 183 adjoined along the coast with Belize, where the larger area of suitable habitat is located, 184 185 however, it is highly unlikely that D. mawii is found in Honduras. The high elevation, mountainous terrain of the Cordillera de Celaque range that begins at the border of Honduras and 186 Guatemala likely forms a strong barrier to movement. Similarly, the Sierra de Chiapas separates 187 the larger contiguous area of high habitat suitability on the Atlantic side of the countries and the 188 isolated area of predicted suitable habitat on the Pacific coasts of Mexico and Guatemala, so it is 189 unlikely that turtles inhabit this region due to the distant, disjointed locations separated by a 190 mountain range. 191

Although MaxEnt is robust against small sample size (Hernandez et al., 2006; Anderson and 192 193 Gonzalez, 2011; Guillera-Arroita et al., 2014) and appropriate subsampling techniques were applied to reduce sampling bias, our presence data includes a potentially biased, small sample 194 size. Due to the fully aquatic lifestyle and small population size (Briggs-Gonzalez et al., 2019), 195 196 locating individuals is difficult and results in a low sample size. Surveys for this species appear 197 to be repeated in some locations and often occurred near villages and cities due to the ease of 198 river access. Because of this, the majority of the points were not evenly distributed throughout the known range but were concentrated in Belize. It is possible that D. mawii is more abundant 199 200 throughout its range than our locality points suggest, furthermore, a more thorough population 201 survey in Guatemala, Mexico, and Honduras may reveal some undocumented sites for D. mawii.

Besides narrowing the potential range for *D. mawii*, the SDM may provide insight for 202 successful reintroduction and protective conservation practices. We found that 31% of highly 203 suitable habitat falls within areas of potentially high hunting pressures. There are no quantifiable 204 data for the long-distance movements of D. mawii in existing literature, yet there are indications 205 that adults move long distances (Briggs-Gonzalez et al., 2019) making it highly likely that 206 207 individuals are traveling through highly suitable areas that are within the human hunting regions (Ellsworth et al. *in prep*). Because overharvesting is the primary driver in local extirpations of D. 208 209 *mawii*, targeting non-fragmented areas, distant from human settlements increases the success of protected areas and reintroduction programs. We cannot be certain that individuals will remain in 210 protected habitats, therefore, the combination of enforcing policy and increased monitoring is 211 even more critical to ensure the survival of this species. 212

We have identified areas that are candidates for protection and release sites, and we have also 213 noted the importance of continued monitoring and enforcement of protective regulations of this 214 215 endangered turtle, but further work is needed. By conducting additional surveys in more remote locations of the species potential distribution in Guatemala, Mexico, and Belize and further 216 research on its ecological niche, the use of protected areas and reintroduction programs can have 217 218 improved outcomes for this and associated species. We can conduct more informed population 219 surveys in areas that have yet to be surveyed to expand our knowledge of finer scale distributions and provide insight to the vulnerability of populations and areas of extirpation. This study also 220 provided environmental associations that create opportunities to compare niches between co-221 222 occurring species and against native predators. Lastly, it is important to investigate other potential range-limiting factors beyond those used in this study to identify other variables 223 affecting this turtle's distribution. 224

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Table 1. Bioclim codes for variables downloaded from the CHELSA database (Kanger et al.,

369 2017). The four variables indicated as important from the PCA (Fig. 2) are bolded.

Code	Variable		
BIO1	Annual Mean Temperature		
BIO2	Mean Diurnal Range		
BIO3	Isothermality		
BIO4	Temperature Seasonality		
BIO5	Max Temperature of Warmest Month		
BIO6	Min Temperature of Coldest Month		
BIO7	Temperature Annual Range		
BIO8	Mean Temperature of Wettest Quarter		
BIO9	Mean Temperature of Driest Quarter		
BIO10	Mean Temperature of Warmest Quarter		
BIO11	Mean Temperature of Coldest Quarter		
BIO12	Annual Precipitation		

Variable		
Precipitation of Wettest Month		
Precipitation of Driest Month		
Precipitation Seasonality		
Precipitation of Wettest Quarter		
Precipitation of Driest Quarter		
Precipitation of Warmest Quarter		
Precipitation of Coldest Quarter		

Table 2. Pearson correlation values for environmental variables. The four variables used for the

373 model are bolded.

Variables	Precipitation of Wettest Month	Precipitation Seasonality	Precipitation of Wettest Quarter	Precipitation of Warmest Quarter	Elevation
Precipitation of Wettest Month	1.0000	-0.0655	0.9951	0.3787	-0.2051
Precipitation Seasonality		1.0000	-0.0725	0.5265	0.3155
Precipitation of Wettest Quarter			1.0000	0.3756	-0.2039
Precipitation of Warmest Quarter				1.0000	-0.1626
Elevation					1.0000

Table 3. Permutation importance of environmental and geographic variables used in SDM

376 model.

Variable	Percent permutation Importance
Elevation	57.2%
Precipitation Seasonality	27.3%
Precipitation of the Wettest Month	8.8%
Precipitation of the Warmest Quarter	6.7%

377



Figure 1. The approximate range of *Dermatemys mawii* in Central America, based on published

381 surveys (Vogt et al., 2011).



Figure 2. A Principal Components Analysis (PCA) of all Bioclim environmental variables at
presence points with the top two highest contributing variables in each principal component
bolded.



Figure 3. Response curves for each raw predictor variable used in the species distribution model.
Predicted value (y-axis) is the predicted habitat suitability from zero (low predicted suitability) to
one (high predicted suitability).


Figure 4. A habitat suitability map of *Dermatemys mawii* ranging from lowest habitat suitability
(0.21) to highest suitability (0.97).





Figure 5. High habitat suitability based on environmental conditions overlain with areas of high
human hunting pressure. High environmental suitability is indicated by the solid grey areas, with
buffered human settlements indicated by the shaded areas. Presence points are included. Large
areas of lower hunting pressures within the highly suitable areas are circled.

409 Chapter 2: Home Range of Juvenile Central American River Turtles

410 (Dermatemys mawii) in Belize

411 ABSTRACT

As anthropogenic factors are leading to species' population declines across the globe, many 412 conservation organizations are implementing initiatives to preserve imperiled species. The 413 Central American river turtle (Dermatemys mawii) is one such imperiled species that has been 414 415 extirpated from areas of its range in Mexico, Belize, and Guatemala. Although this culturally 416 important species has long been harvested for sustenance, more recent unsustainable hunting has had a negative impact on populations. Home range information is crucial to conservation 417 418 initiatives that protect critical habitats and headstart programs. Here, we tracked movements of juveniles at two sites: one with substantial human hunting activity and one protected site in a 419 river in Belize. We found that juveniles maintain relatively small home ranges with a mean of 420 421 3.4 ha, but occasionally move out of these ranges. Despite the difference in hunting pressure, 422 there were no difference in movement distances or home range size between the two sites. Juvenile turtles had a higher capture rate relative to adults at the human activity site, suggesting 423 that human presence may reduce natural predators of juveniles, but such an advantage is likely 424 lost as turtles grow and become targets of hunting. Future studies assessing the effects of human 425 426 activity and overexploitation on the movement and survival of *D. mawii* are crucial in aiding conservation initiatives. 427

428 INTRODUCTION

Habitat loss, habitat fragmentation (Fahrig, 1997; Marchand and Litvaitis, 2004; Böhm et al.,
2016), climate change (Thomas et al., 2004; Walther et al., 2002; Wiens, 2016), and

overexploitation (Klemens and Thorbjarnarson, 1995; Milner-Gulland and Bennett, 2003; 431 Allendorf et al., 2008; Ripple et al., 2019) are causing devastating population declines, putting 432 species at risk of extinction, and consequently impacting ecosystems. Turtle and tortoise 433 populations, in particular, have been impacted by landscape modification and overexploitation 434 causing nearly two-thirds of these species to be classified under a heightened risk of extinction 435 436 by the International Union for Conservation of Nature (IUCN Red List). Habitat alteration from urbanization and agricultural development lead to a decrease in suitable nesting sites (Marchand 437 438 and Litvaitis, 2004), while increasing both the risk of vehicle collisions (Gibbs and Shriver, 439 2002; Bowne et al., 2006; Dorland et al., 2014) and predation from species associated with urban and suburban development (Seigel, 1980; Congdon, 1993; Marchand and Litvaitis, 2004). 440 Similarly, over-harvesting of breeding populations for sustenance or commercial trade negatively 441 affects the viability and health of the remaining populations (Klemens, 2000; Zimmer-Shaffer et 442 al., 2014). Despite severe anthropogenic threats, freshwater turtles remain underrepresented in 443 444 conservation research (Burke and Gibbons, 1995; Roll et al., 2017; Lovich et al., 2018), mirroring an overall gap in conservation of freshwater ecosystems (Abell, 2002) and further 445 complicating the ability to conserve freshwater turtles. 446

Combining knowledge of life history and environmental factors with field studies is critical to improve conservation efforts (Wilson et al., 2006; Griffith et al., 1989; Fischer and Lindenmayer, 2000; Brooks et al., 2006; Berger-TAL and Saltz, 2014; Yang et al., 2018). For example, habitat loss and overexploitation can be opposed with long-term conservation programs that offer protection from human-induced mortality and assist in the breeding success of target species; thus, these programs are fundamental to preserving biodiversity and the survival of threatened turtle species (Ceballos and Ehrlich, 2010). Such conservation initiatives have a higher

likelihood of success when key components of a species' ecology, home range, and movement 454 patterns are incorporated into conservation strategies (Burt, 1943; Hart, 1983; Gibbons et al., 455 456 1990; Powell, 2000, 2012; Lidgard et al., 2020). Spatial studies provide insight into population distribution, dispersal patterns, and habitat selection (Bowler and Benton, 2005), which inform 457 potential release sites and allow for assessment of reintroduction success for programs aimed 458 459 toward augmentation and reintroduction (Fischer and Lindenmayer, 2000; Cadi and Miquet, 2004; Berger-TAL and Saltz, 2014). Home range studies provide minimal habitat requirements 460 that can also be used to designate protected areas and ensure adequate space for individuals to 461 carry out their activities with minimal human impact (Allen and Singh, 2016; Di Franco et al., 462 2018); moreover, species' preservation in protected areas has a higher success rate when the size 463 of the protected habitat is considerably larger than the species home range size (Di Franco et al., 464 2018; Green et al., 2015). 465

The Central American river turtle (*Dermatemys mawii*) is listed as Critically Endangered by 466 the IUCN (Vogt et al., 2006). This freshwater turtle is found primarily in lowland rivers, lakes, 467 lagoons, and brackish waters in the Yucatan peninsula region of Central America (Iverson and 468 Mittermeier, 1980; Legler and Vogt, 2013; Briggs-Gonzalez et al., 2019). The primary threat to 469 470 D. mawii is overexploitation, which has extirpated and fragmented populations across its range 471 (Iverson and Mittermeier, 1980; Alvarez del Toro et al., 1979; Vogt et al., 2011). In other turtles 472 and aquatic vertebrates, pressure from overhunting has caused population declines through decreased clutch sizes (Daza and Páez, 2007; Eisemberg et al., 2017), reduced recruitment 473 474 (Marchand and Litvaitis, 2004; Walsh et al., 2006), and restricted genetic variability (Ratner and Lande, 2000; Walsh et al., 2006; Allendorf et al., 2008). Similar to other turtles, D. mawii has 475 slow population growth because of low offspring survivorship and long maturation times, 476

making populations particularly vulnerable to over-harvesting (Dunham et al., 1989; Congdon,
1993; Gibbons et al., 2000). Although the exact time of reproductive maturity and longevity in
the wild is undocumented, the generation time for this species is thought to be approximately 10
years (Briggs-Gonzalez et al., 2019) and life expectancy exceeds 30 years in captivity (BriggsGonzalez et al., 2019).

To counter the serious threats *D. mawii* is facing, spatial studies specifically investigating 482 483 home range size are needed to better understanding habitat use in order to inform conservation initiatives that protect and bolster existing populations. The bulk of information on D. mawii has 484 investigated the species' population size (Rainwater et al., 2012), diet (Moll, 1989), population 485 486 genetics (González-Porter et al., 2011, 2013), habitat preference (Legler and Vogt, 2013), and conservation threats (Rangel-Mendoza et al., 2009, 2014; Polisar and Horwich, 1994). However, 487 much of the current spatial information on D. mawii stems from published observations and 488 personal communications with people living near or within the species' range (Alvarez del Toro 489 490 et al., 1979; Iverson and Mittermeier, 1980; Moll, 1989; Polisar, 1992, 1996; Rainwater et al., 2012; Vogt et al., 2011; Legler and Vogt, 2013). These studies, personal communications, and 491 anecdotes do not quantify movements, but suggest that adults undergo long-distance movements 492 493 and offer hypotheses about factors that may influence movement patterns and home range size, 494 including seasonal effects. The rainy season, especially, plays an important role in shaping home 495 ranges and movement patterns of adult D. mawii (Briggs-Gonzalez et al., 2019; García-Anleu et al., 2010; Legler and Vogt, 2013). Specifically, turtle dispersal activity increases during the rainy 496 497 season when floodplains and flooded forests become inundated (García-Anleu et al., 2010; Legler and Vogt, 2013). Conversely, during the dry season, the reduced rainfall and subsequent 498 low water levels limit the amount of available habitat. Low water levels force turtles to 499

aggregate, providing a prime opportunity to encounter other individuals, and eliminating the
need to travel long distances to search for a mate (Polisar, 1992, 1996; Briggs-Gonzalez et al.,
2019). Consequently, receding water levels and aggregated populations increase vulnerability to
human hunting. Although some factors that affect the movements of adult *D. mawii* are known, it
remains unclear whether juveniles follow the same home range attributes and seasonal
movement patterns as the adults.

506 Home range and movements of many animals are driven by a combination of intrinsic 507 biological and extrinsic environmental factors that may vary among seasons or over an individual's lifetime. For example, reproductive maturity can influence home-range attributes 508 509 and result in disparities in home range size between juveniles and adults (Gibbons et al., 1990; McNab, 1963; Beck-King et al., 1999; Sutherland et al., 2000). Reproductively mature 510 individuals of many species often expand their home range and increase movement to optimize 511 reproductive success (Morreale et al., 1984; Doody et al., 2002; Slavenko et al., 2016), whereas 512 513 reproductively immature individuals benefit from maintaining a smaller home range and shortrange movements, which can reduce predation (Morreale at al., 1984). Juvenile turtles heavily 514 515 invest into growth and development of the protective shell rather than overall size, leaving them 516 at a disadvantage to escape predation (Moll, 1986; Alvarez del Toro, 1982; Smith and Smith, 517 1980; Platt et al., 2016). Naturally high juvenile predation, in combination with increased 518 anthropogenic hunting, affect both ends of the age spectrum and have potential impacts on 519 population stability (Moll and Moll, 2004). Although not always the case (Crouse and Frazer, 520 1995; Sung et al., 2013), high juvenile representation often implies stable turtle populations. 521 Although survivorship rates have not been documented for *D. mawii*, populations would still benefit from conservation efforts that protect juveniles. 522

Here, we aim to quantify home ranges and describe movement patterns of juvenile *D. mawii* from two locations in central Belize using VHF (Very High Frequency) transmitters. Given that juveniles are more susceptible to predation and lack the need to search for a mate, we expect juveniles to move shorter distances and occupy a smaller area than adults. Investigating the movement and home range size of juvenile *D. mawii* will provide critical data to improve conservation efforts, inform policy makers, and increase our understanding of the species' ecology.

530 MATERIALS AND METHODS

531 Study site

532 We conducted our study at two locations, during two different dry seasons in a large, slow 533 moving, perennial stream that flows through lowland tropical savannah into a series of lakes. 534 Over the period of study, air temperatures ranged from 23°C–29°C with a total of 4.6 cm of 535 rainfall (2019) and 23°C-31°C with 2.0 cm of rainfall (2020). Maximum water depth was 5 m. The riparian vegetation at these sites is dominated by emergent plants such as *Mimosa pudica* 536 537 and Paspalum paniculatum. The prevailing trees, Manilkara bidentata, and Haematoxylum sp., provide canopy cover along the bank and contribute to the coarse woody debris found throughout 538 the stream. We chose the first site (A) based on accessibility and quality of habitat, then moved 539 540 to the second site (B), 10 km south of site A and within a wildlife sanctuary, due to the high presence of human interference at site A. Site A, which was publicly accessible, had a high 541 occurrence of canoeing, net trapping, and poaching activity, whereas site B had minimal 542 anthropogenic activity. We tagged individuals at site A between January and March in 2019 and 543 between February and March 2020 at site B. 544

545 Sampling

In both 2019 and 2020, river accessibility limited our study to the dry season. Individuals of D. 546 mawii were captured and tagged at the beginning of the dry season between January and March, 547 just as daily precipitation declined, and water levels began to recede (Lambert et al., 1980). 548 Trammel nets (45 m long x 3 m deep, outer wall mesh size of 355 mm x 355 mm, inner wall 549 mesh size of 101 mm x 101 mm) were left open along banks or across the stream by tying the 550 551 ends to branches of overhanging trees, submerged logs, or available sturdy woody stems along the bank during peak turtle activity time, largely between 1800 h and 0500 h (Rainwater et al., 552 553 2012; Legler and Vogt, 2013) for 17 net nights at site A and 16 net nights at site B. Net checks 554 were conducted every 90 minutes to ensure the safety of the turtles; no turtles drowned during the course of this study. All turtles, except one small hatchling that was captured by hand, were 555 captured using the trammel net. The large mesh opening precluded juveniles under 144 mm. For 556 each individual, we recorded the mass using a spring scale (Pesola) accurate to the nearest gram. 557 We recorded the straight-line carapace and plastron lengths using manual calipers to the nearest 558 1.0 mm. 559

Waterproof VHF transmitters (RI-2B, Holohil) were attached using an epoxy glue (Devcon) to 560 the fourth costal scute of the carapace to minimize any impacts on foraging, predator evasion, 561 562 and entanglement with vegetation. Transmitters had a pulse rate of 40 ppm, a pulse width of 24 563 ms, and a total weight of 15 g. In addition to the VHF transmitter, each individual was marked 564 with a prepackaged sterile passive integrated transponder (PIT) tag (GPT12, Biomark) with a 565 unique identification number. The PIT tag was inserted sub-dermally in the right hind limb of 566 each individual using sterile techniques. This tag allowed turtle identification in the event of 567 transmitter detachment. Individuals with a carapace length of greater than 365 mm were classified as adults, and smaller individuals were classified as juveniles (Legler and Vogt, 2013). 568

The sex of each adult was determined based on the presence or absence of sexually dimorphic 569 characteristics. At sexual maturity, males display external, visual characteristics such as bright 570 571 yellow coloration of the head and a tail extending beyond the carapace, whereas females lack the brightly colored head and long tails (Legler and Vogt, 2013). Juveniles could not be visually 572 sexed due to lack of sexual characteristics. No bias towards age or sex was intentional; all 573 574 individuals captured were tagged and used for this study. Duration of transmitter attachment was limited by annual molting of the outer, keratinized layer of scutes, which often occurs in the late 575 576 dry season for *D. mawii* (Legler and Vogt, 2013). No individuals had visual signs of molting at 577 the time of capture. Animals were held for a maximum of 60 minutes to provide sufficient time for the epoxy glue to set and cure prior to release at the site of capture. All appropriate research 578 permits were obtained (MSR permit 0015-20) and the study was approved by the University of 579 Oklahoma's Institutional Animal Care and Use Committee (R19-002, R19-002A). 580

581 Radio telemetry, data collection, and distance measurements

After we tagged and released turtles, they were tracked manually from a canoe using a handheld radio receiver (R-1000 Telemetry Receiver, Titley Scientific, Columbia, MO) and a 3-element Yagi antenna (Titley Scientific, Columbia, MO). For both sampling periods, tracking occurred between 2 to 7 times per week following transmitter attachment. Individuals at site A were tracked for five consecutive weeks (January–March 2019) and at site B, individuals were tracked three weeks (February–March 2020).

588 Location data were predominately recorded during peak activity hours because of the species'

tendency to rest at the bottom of riverbed when not active (Briggs-Gonzalez et al., 2019) making

them difficult to locate. We occasionally tracked individuals at different times in a 24-hour

591 period to investigate movement activity at various times of the day. Our tracking area covered

over 500 meters both up- and downstream from the initial capture sites for a total of about 3 km 592 stretch of river. When an individual was located, we recorded the time of day and used a GPS 593 (GPSMAP 010-01199-10, Garmin, WGS84) to record coordinates of its location (or fix). All 594 distances were calculated using the Euclidean (straight-line) distance formula using the "dist" 595 function in the base R package. The sum of the distances between each locality fix was divided 596 597 by total number of fixes to calculate the mean distance per fix (mean displacement distance) traveled by each individual. The mean displacement distance was then estimated for the mean 598 599 number of days between detections. Although measures were taken to prevent disturbing the individuals from their location, if a turtle fled in response to being approached, GPS coordinates 600 were recorded after following individuals for five meters. 601

602 Home range calculations and statistical analyses

603 To estimate home range, we used 100% and 95% minimum convex polygons (mcp), which 604 provides a more accurate estimation of home range for small sample sizes (Boyle et al., 2009). 605 MCPs create a perimeter representing the minimum home range of each individual around the five outermost GPS coordinates, so that it encompasses these points and all other points where an 606 607 individual was located using either 100% of the locality points, or the points that fall in the 95% 608 quantile. We then buffered the perimeter of each home range polygon by 15 m to account for any variation in GPS signal, following the methods of Ryan et al. (2008). The area of each mcp was 609 calculated in the adehabitatHR package (Calenge, 2006) in R (R Core Team, 2019, Version 610 611 1.2.5019) and adjusted to fit along the river. We conducted a series of correlation tests using the base cor.test function in R to detect relationships between carapace size, number of fixes, and 612 613 home range size. Spearman rank correlations were used to test the effect of size using carapace length on the number of fixes we were able to obtain and on the association between turtle size 614

and home range size. A Spearman rank correlation was also used to determine whether the number of GPS location points affected home range size. Due to inconsistencies in the tracking effort, detection, and duration, we were unable to statistically compare home range or movement between sites. We used a Fisher's exact test to determine if there was a difference in our ability to locate juveniles or adults. We used $\alpha = 0.05$ as the statistically significant level for all tests.

620 **RESULTS**

621 Sampling and detectability

622 Seventeen juveniles and four adults were captured via trammel net or hand. Thirteen juveniles and one adult were captured at site A over 17 net nights with an average of nine fixes per 623 624 individual over five weeks. Five juveniles and three adults were captured at site B over 16 net 625 nights with an average of five fixes over three weeks. On average, individuals were detectable 626 (i.e., had greater than one detection) at site A nearly every two days (47% of the attempted tracking time), while individuals at site B were detectable approximately every four days (35% 627 of the attempted tracking time). Home range estimates and mean displacement distances were 628 629 calculated for a total of ten individuals that had a minimum of five locality fixes (Table 1) at site A (Fig. 1) or site B (Figs. 2 & 3). 630

Tracking and trapping times indicate that individuals were more likely to be located between
2000h and 0600h, which coincides with their previously recorded peak activity times (BriggsGonzalez et al., 2019). However, some individuals were active and relocatable during daylight
hours (0600 – 2000 h). The majority of the individuals captured were juveniles, with a smaller
proportion of adults to juveniles at site A (0.375) than site B (0.077). Three out of the four adults
captured for this study were re-detected after initial capture (Fig. 4).

637 Home range size

Juvenile home range size (95% mcp) varied from 0.27 ha to 2.14 ha at site A (Fig. 5a) and 0.48 638 ha to 21.14 ha at site B (Fig. 5b). Mean home range size (95% mcp) appears larger at site B (6.5 639 ha) than at site A (0.8 ha). One-half of the individuals tracked included one or two locality points 640 deviating from the core occupied area. We found that turtle size (carapace length) did not affect 641 the home range size (S = 234, p = 0.2324; Fig. 6a), nor the number of locality fixes (S = 174.46, 642 643 p = 0.2198; Fig. 6b). The number of fixes varied among individuals, but a greater number of locality detections was not correlated with a larger home range area (S = 112.68, p = 0.372; Fig. 644 7). 645

646 Movement

647 Mean displacement movement between points of the nine tracked juveniles resulted in an

average movement of 127.4 meters per every two days at site A and an average movement of

649 277.1 meters every four days at site B. Turtle size had no effect on mean displacement distance

650 (S = 372, p = 0.9432; Fig. 6c), nor likelihood of detection (p = 0.5865).

The number of fixes were below the minimum required to calculate home range for all but one adult (EDE 232). EDE 264 traveled 1780 m north from the initial capture site in less than 24 hours and was not located again within the 2 km of its original capture in the remainder of the study. The other adult, EDE 262, was not located within the search area (2 km from the original capture location) for a month before reappearing 277 m from its original net capture location.

656 **DISCUSSION**

Juveniles had relatively small home ranges and typically remained near the area where theywere initially captured, even when multiple individuals were captured in the same net. Our

results also indicate that juveniles occasionally travel outside a core area. These findings indicate that allocating protected lands may be sufficient to benefit juveniles of this species during the dry season. Both capture and detection rates for adults in our study were too low to allow direct comparison of home range sizes between juveniles and adults; however, observations of adults by local community members and previous studies indicate that adults make very long-distance movements (Alvarez del Toro et al., 1979; Iverson and Mittermeier, 1980; Moll, 1989; Polisar, 1992, 1996; Rainwater et al., 2012; Legler and Vogt, 2013).

666 Our observations and inability to reliably track adults indicate that they are either occupying 667 extensive home ranges beyond our survey area or they make far-ranging movements that 668 provided us the opportunity to capture them while they traveled through our study site. However, 669 the latter can be detrimental if the adult is moving through an area with high human hunting 670 activity. These observations coupled with our data on juvenile movement also indicate a 671 potential ontogenetic change in the home range of this species.

672 Human disturbance by frequent boating activity, as occurs at site A, can impact nesting and basking behavior (Moore and Seigel, 2006), stress levels, and overall health of individuals 673 674 (Selman et al., 2013). We suspect human activity plays a large role in the movements and home ranges of these individuals based on studies of similar lotic species and from personal 675 observations. Although we could not directly compare home range size and mean fix distances 676 between sites, long-term data and a larger sample size may show that human activity and 677 678 disturbance result in smaller home range sizes and shorter movements. We observed longer distances and greater home range size at site B which could likely be a consequence of the 679 680 difference in human activity between sites. At site B, hunting and fishing are regulated by local personnel and very little boat activity was observed. Hunting activity could also be directly 681

responsible for the lack of adult captures. Interestingly, the human activity at site A could also be suppressing the juveniles' natural predators, such as crocodiles, thereby decreasing juvenile mortality rates and producing the observed higher capture rate of juveniles relative to adults at sire A. Any indirect protection of juveniles by hunting and other human activity would be lost as turtles grow and become targets for human sustenance.

Only one-half of individuals that we captured for this study had the minimum number of fixes 687 688 to be used for the mcp calculations, possibly due to VHF limitations and the life history of D. *mawii*. Although VHF technology is useful in pinpointing an individual, the signal attenuates 689 with depth or at great distances, which is likely the reason for few data points for adults. Other 690 691 technologies, such as GPS and satellite technologies, provide more spatial information with higher accuracy, but both technologies are aerial systems that require an individual to breach the 692 surface to send a signal. As this species is wholly aquatic and does not leave the water to bask. 693 Based on current available technology, VHF provides better insight into fully aquatic freshwater 694 695 species that live in submerged holes along the bank (Skupien et al. 2016), however, testing a combination of technologies that transmit a strong signal from greater distances, such as 696 ultrasonic transmission or acoustic tracking systems, is needed to determine the optimal tracking 697 698 method for this fully aquatic organism.

Our study provides quantitative data on the movements and home ranges of these individuals, now identifiable with PIT tags, for future studies. With increased tracking duration and improved technology, we can better understand the home ranges of *D. mawii* at various life stages and across seasons. Our study was limited to the dry season when sites are most assessable, but seasonality is a driver for movement in other freshwater turtles (Litzgus and Mousseau, 2004; Fachín-Terán et al., 2006) and is therefore imperative to assess home ranges across seasons and

- across years. Our study also suggests the difference in human presence potentially plays a role in
- movement and home range of individuals, however extensive studies are needed to better
- 707 understand how anthropogenic activity affects this species.

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ID number	Age/ Sex	100% mcp (ha)	95% mcp (ha)	Number of Fixes	Maximum Distance Between Fixes (m)	Mean Displacement Distance (m)	Carapace length (cm)			
Site A										
223*	Juvenile/ Unknown	3.830	2.14	21	608.90	175.44	28.0			
224*	Juvenile/ Unknown	0.510	0.37	13	71.42	29.55	27.0			
225*	Juvenile/ Unknown	2.520	0.87	8	522.59	197.29	28.2			
226	Juvenile/ Unknown	NA	NA	1	NA	NA	30.1			
227	Juvenile/ Unknown	NA	NA	1	NA	NA	27.1			
228	Juvenile/ Unknown	NA	NA	1	NA	NA	35.4			
229	Juvenile/ Unknown	NA	NA	1	NA	NA	34.2			
230	Juvenile/ Unknown	NA	NA	1	NA	NA	33.1			
231*	Juvenile/ Unknown	0.420	0.41	12	512.26	132.63	30.5			
233	Juvenile/ Unknown	NA	NA	1	NA	NA	35.2			
234*	Juvenile/ Unknown	0.950	0.27	5	270.61	101.84	27.5			
235	Juvenile/ Unknown	NA	NA	1	NA	NA	17.4			
232*	Adult/ Female	0.260	0.15	6	47.17	27.36	41.5			

Table 1. Movement and home range data for each relocated individual.

ID number	Age/ Sex	100% mcp (ha)	95% mcp (ha)	Number of Fixes	Maximum Distance Between Fixes (m)	Mean Displacement Distance (m)	Carapace length (cm)
Site	В						
259*	Juvenile/ Unknown	1.930	1.39	8	341.04	225.00	26.1
260*	Juvenile/ Unknown	55.220	3.08	15	6,604.82	915.33	24.2
261*	Juvenile/ Unknown	28.469	21.14	6	3,522.11	811.46	28.7
263	Juvenile/ Unknown	NA	NA	2	30.08	30.08	29.3
265*	Juvenile/ Unknown	0.700	0.48	7	90.88	56.12	26.5
258	Adult/ Female	NA	NA	1	NA	NA	40.9
262	Adult/ Female	NA	NA	2	277.06	277.06	42.3
264	Adult/ Female	NA	NA	2	1,779.84	1,779.84	37.8
*Individuals used in mcp calculations							







Figure 2. 100% minimum convex polygon (mcp) home range estimates for four juveniles at site
B represented using unique colors. Each point represents an individual fix, and the surrounding
color-shaded polygon shows the 100% mcp home range estimates.



Figure 3. 95% minimum convex polygon (mcp) home range estimates for four juveniles at site B
represented using unique colors. The grey band shows the 95% mcp home range of each
individual. Flow direction is indicated by the arrow.



Figure 4. A map at site B indicating the three individuals that were only relocated a single time
following initial capture. Initial capture is indicated by the 'i' adjacent to the locality fix. Stream
flow is indicated with an arrow.



949 Figure 5. Variation in 95% mcp home range sizes at sites A (a) and B (b).



(a)







Figure 7. The number of fixes of each individual with greater than 5 fixes does not predict homerange size.