



Testing the efficiency of protected areas in the Amazon for conserving freshwater turtles

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ABSTRACT

Aim We used chelonian distribution data to (1) predict suitable areas of the occurrence for freshwater turtle species using species distribution models and (2) evaluate whether these species are protected by the current network of protected areas (PAs).

Location The Brazilian Amazon.

Methods We generated predictions of suitable areas for chelonian occurrence based on BIOCLIM, SVM, GLM and maximum entropy modelling procedures. We used maximum entropy to run the gap analysis and compared the effectiveness of three kinds of protected areas with different levels of protection: (1) integral protection areas (IPA) only; (2) integral protection areas + sustainable use areas (IPA+SUA); and (3) integral protection areas + sustainable use areas + indigenous lands (IPA + SUA + IL).

Results We identified only one full gap species, *Mesoclemmys nasuta*, whose distribution is not included in any PAs. Other chelonian species have at least a portion of their distribution included in PAs. Some protected species and partial gap species occur in areas with high rates of deforestation. Considering PAs with the highest level of protection (IPA), only *Rhinoclemmys punctularia* and *Kinosternon scorpioides* achieve their conservation targets. In the IPA + SUA scenario, conservation targets of some species with small range sizes are not achieved. When all PA types were considered (IPA + SUA + IL), only two species fail to achieve their conservation targets, *Acanthochelys macrocephala* and *M. nasuta*.

Main conclusions Despite the large number of PAs in the Brazilian Amazon, IPAs alone are not sufficient for capturing suitable areas for freshwater turtles. The inclusion of SUA and IL is crucial for achieving coverage targets for most species. However, chelonians may be overharvested in SUAs and ILs, due to their importance as a food resource. Areas that have high turtle richness next to existing PAs and the needs of traditional cultures should be considered in management planning for freshwater turtles.

Keywords

Amazon, gap analysis, turtle conservation, vulnerability of freshwater organisms.

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INTRODUCTION

The need for conservation planning is particularly urgent in the tropics (Klink & Machado, 2005; Cayuela *et al.*, 2009) where habitat loss and degradation contribute to the decline in fauna, generating what is known as the ‘biodiversity crisis’

(Myers, 1996). For instance, most megadiverse areas currently occur in the tropics (Myers *et al.*, 2000) and the Amazon includes ecoregions with high levels of richness and endemism of aquatic organisms (Abell *et al.*, 2008). Deforestation in the Amazon basin is driven by socio-economic development, mainly cattle ranching (Fearnside, 2005, 2008;

Macedo *et al.*, 2012; Castello *et al.*, 2013; Souza *et al.*, 2013). A large proportion of the basin has been deforested or altered, and deforestation rates since 1991 have trended upward (Fearnside, 2005). To decrease threats associated with deforestation, it has been suggested that megareserves be created to represent different biological assemblages, including aquatic vertebrates (Peres & Terborgh, 1995; Peres, 2005).

Knowledge about species' distributions is an important basic piece of information for conservation planning and prioritization (Peres, 2005; Thieme *et al.*, 2007). Lack of information about biogeography and the distribution of organisms, the so-called Wallacean shortfall (Lomolino, 2004; Diniz *et al.*, 2010), is widely recognized as a critical limitation for effective management actions, especially in tropical regions (Myers *et al.*, 2000; Brooks *et al.*, 2001). Frequently the only available information about species distributions is range maps, which are typically coarse overestimates of species occurrence (Rodrigues *et al.*, 2003; Rondinini *et al.*, 2006; Hurlbert & Jetz, 2007). Records for most chelonian species in the Amazon are limited to a few localities within their ranges (Souza, 2004, 2005; Brito *et al.*, 2012). In this context, predictive distribution models can be an important tool to fill gaps in knowledge about species' distributions (Raxworthy *et al.*, 2003; Costa *et al.*, 2010). These models are commonly called species distribution models (SDMs) (Araújo & Peterson, 2012; Peterson & Soberón, 2012) specially in studies that try to generate hypotheses about species distributions, rather than modelling their niche (Van Loon *et al.*, 2011).

Independent of the terminologies that are used, predictive distribution models have the same purpose, to identify suitable habitat for populations of a species (Guisan & Thuiller, 2005; Elith & Leathwick, 2009; Franklin, 2010; Peterson *et al.*, 2011), through identification of statistical relationships between species' occurrences and a set of environmental predictors (Guisan & Zimmermann, 2000). Suitable areas can be then projected into geographic space to estimate species' geographic distribution (Peterson, 2001). These analyses are performed using different modelling procedures, depending on different theoretical conditions and assumptions (Elith *et al.*, 2006; Austin, 2007; Elith & Leathwick, 2009). Different methods often show substantial variation in performance (Elith *et al.*, 2006; Peterson *et al.*, 2007).

Species distribution models are useful for management (Peterson *et al.*, 2001; Guisan & Thuiller, 2005; Araújo *et al.*, 2011; Crowder & Heppell, 2011; Nóbrega & De Marco, 2011) because they produce maps showing the environmental suitability for species occurrence in areas that have not been previously sampled and can produce valuable information about overall spatial patterns in biological diversity (Cayuela *et al.*, 2009; Nóbrega & De Marco, 2011). Thus, these models are advantageous for evaluating the efficiency of existing protected area (PA) networks in representing species distribution, as assessed in formal gap analyses (Rodrigues, 2003; Phillips *et al.*, 2006; Loucks *et al.*, 2008).

Protected areas have been an effective tool for maintaining viable populations of threatened species or species potentially impacted by human occupation (Rodrigues *et al.*, 2003; Sánchez-Azofeifa *et al.*, 2003; Veríssimo *et al.*, 2011). However, gap analyses have demonstrated that existing PA networks in the Americas are usually inadequate to conserve biodiversity (Scott *et al.*, 2001; Ochoa-Ochoa *et al.*, 2007).

The applicability of SDMs in the freshwater aquatic realm has been poorly explored (Wiley *et al.*, 2003) due to the lack of distribution data for freshwater species (Thieme *et al.*, 2007) and limited data describing local environmental conditions (Iguchi *et al.*, 2004; McNyset, 2005; Oakes *et al.*, 2005). Freshwater biodiversity has been more impacted than the most of terrestrial organisms (Sala *et al.*, 2000). However, priority areas for conservation are typically established based on terrestrial species and ecosystems (Brooks *et al.*, 2006; Castello *et al.*, 2013), and aquatic habitats are only protected by chance (Skelton *et al.*, 1995; Peres, 2005). Conservation planning and strategies that encompass both terrestrial and aquatic environments are crucial for effective management, especially in Amazon, where freshwater ecosystems cover between 14 and 29% of the basin area (Thieme *et al.*, 2007; Castello *et al.*, 2013).

Turtles are one of the most endangered groups of vertebrates (Gibbons *et al.*, 2000; Van Dijk *et al.*, 2000; Turtle Conservation Fund 2002; IUCN, 2011). Böhm *et al.* (2013) estimated that 52% of freshwater turtles are threatened. Of the 16 freshwater species of turtles in the Brazilian Amazon, seven are in some threat category (IUCN, 2011). In this context, the knowledge about current distribution patterns of turtles and the contribution of PAs to their conservation could not be more important (Iverson, 1992a; Stuart & Thorbjarnarson, 2003; Rhodin, 2006). Thus, our objectives in this study are to (1) predict suitable areas of occurrence for freshwater Amazon chelonians and (2) evaluate whether the group is protected by the existing network of Amazonian PAs.

METHODS

Species occurrence records

We compiled an occurrence database for 16 freshwater turtles (see Table 1) including data from the following sources: an extensive literature review, Brazilian scientific collections and museum specimens obtained from Species Link (CRIA, 2015), unpublished data from our research group and from a governmental project, Projeto Quelônios da Amazônia (IBAMA, 2015a). In addition, we utilized species data provided by the EMYSsystem Global Turtle Database (Iverson *et al.*, 2003), which records depict the maps produced by Iverson (1992a,b,c). To minimize modelling problems caused by errors in georeferencing, we deleted occurrence records that were obviously erroneous, records with imprecise geographic coordinates, and generalized location descriptions. This process resulted in 1826 occurrence records (Table 1).

Table 1 The number of spatially unique occurrence records (at 4 km² resolution) for 16 freshwater turtles in Brazilian Amazon. We also show the amount of suitable habitats (km²), the proportion of the conservation targets (%) and the proportion of the conservation targets attained (%) for those species using (i) only the integral protection areas (IPA), (ii) integral protection areas + sustainable use areas (IPA + SUA) and (iii) integral protection areas + sustainable use areas + indigenous lands (ITA + SUA + IL).

Species	Unique records	Suitable habitats	Conservation target	IPA	IPA + SUA	IPA + SUA + IL
Semi-aquatic						
<i>Kinosternon scorpioides</i>	67	2,915,552	10	10.7	27.3	45.1
<i>Rhinoclemmys punctularia</i>	40	1,602,432	10	11.3	21.2	44.2
<i>Acanthochelys macrocephala</i>	13	91,360	50.5	19.3	25.4	40.5
<i>Mesoclemmys vanderhaegei</i>	18	222,864	35.9	9.9	23.8	43.4
<i>Mesoclemmys gibba</i>	48	4,111,632	10	6.4	15.6	29.3
<i>Platemys platycephala</i>	45	2,281,552	10	7.1	12.9	27.7
Aquatic						
<i>Chelus fimbriata</i>	71	1,676,768	10	5.5	22.5	34.1
<i>Mesoclemmys raniceps</i>	28	3,489,664	10	7.7	22.9	39.6
<i>Mesoclemmys nasuta</i>	11	10,336	81.7	0	0	0.07
<i>Phrynops geoffroanus</i>	39	1,799,584	10	5.8	11.9	29.9
<i>Rhinemmys rufipes</i>	13	1,416,640	10	9.1	29.2	42.9
<i>Peltecephalus dumerilianus</i>	78	802,768	10	9.8	28.1	37.6
<i>Podocnemis erythrocephala</i>	97	1,537,360	10	8.7	23.1	35.8
<i>Podocnemis expansa</i>	305	2,147,648	10	7.1	22.1	35.1
<i>Podocnemis sextuberculata</i>	168	2,085,968	10	7.4	22.8	37.1
<i>Podocnemis unifilis</i>	329	2,107,616	10	7.5	22.9	35.5

We included in the analyses not only exclusively aquatic species, but also semi-aquatic species, that live in small temporary and perennial water bodies in forests. As such, we covered the entire area of the Brazilian Amazon in our modelling efforts, as opposed to only including the aquatic ecosystems. The area was divided into a grid of approximately 4 km² cells. We considered only one occurrence record of each species in each cell (spatially unique records) to help avoid effects of sampling bias (Dennis & Thomas, 2000; Kadmon *et al.*, 2004) (Table 1).

Environmental data

Aquatic organisms are influenced by a suite of local environmental variables (Mendonça *et al.*, 2005) for which spatial information is not readily available. However, some studies have shown that macroscale variables performed similarly to local variables when modelling the distribution of aquatic species (Watson & Hillman, 1997; Porter *et al.*, 2000). In the Brazilian Amazon, limnological and macroscale predictors are highly correlated (Frederico *et al.*, 2014). Following this reasoning, we used 42 variables: 37 climatic predictors, three variables that reflect terrain shifts and two predictors that characterize the aquatic environment (see Appendix S1 in Supporting information). We performed a principal components analysis (PCA) of the environmental variables to decrease collinearity among them and to avoid model overfitting. For the PCA, we compiled all layers at a resolution of 4 km². The PCA scores were used as environmental layers in the SDM procedures (Jiménez-Valverde *et al.*, 2011; Dormann *et al.*, 2012). Considering the Kaiser-Guttman criterion of princi-

pal components selection (Peres-Neto *et al.*, 2005), we selected 12 principal components which were responsible for more than 95% of the variation in the environmental variables data (see Appendix S2). We then used these principal components as predictor variables to develop our SDMs (Guisan & Thuiller, 2005; Peterson *et al.*, 2011).

Species distribution modelling

We calculated four different statistical methods for modelling to provide a more reliable estimate of the distribution of turtles (Rocchini *et al.*, 2011): a 'presence-only' method called BIOCLIM (Nix, 1986; Piñero *et al.*, 2007); a 'presence/pseudoabsence' approach via generalized linear modelling (GLM – Stockwell & Peters, 1999; Guisan *et al.*, 2002); and two-class support vector machines (SVM – Schölkopf *et al.*, 2001; Tax & Duin, 2004; Guo *et al.*, 2005). These methods relate known occurrence localities with 'pseudoabsences' extracted from sites at which the species is not known to occur in the study area (Peterson *et al.*, 2011). In addition, we used one 'presence/background' approach, maximum entropy (Phillips *et al.*, 2006; Phillips & Dudik, 2008; Elith *et al.*, 2010). This approach assesses the relation between the environment at the locations of known records and the environment across the entire study area (Peterson *et al.*, 2011). We used the software MaxEnt to run maximum entropy (Phillips *et al.*, 2006), and the 'dismo' package on R Software (R Development Core Team 2012) to run the other modelling methods. Considering possible restriction of accessibility (Barve *et al.*, 2011), we created and evaluated all models for the entire Amazon basin.

We divided occurrence data of species that had more than 15 spatially unique records into 80–20% training–test subsets. We used the training subset to fit the SDMs and the test subset to evaluate the predictions. We based the evaluation of model performance on the elements of a confusion matrix or on the measures derived from this matrix (Elith *et al.*, 2006; Peterson *et al.*, 2011). We used 10,000 random pseudoabsence localizations for GLM and SVM methods and 10,000 background data for maximum entropy. For species that had < 15 spatially unique records, we fit and tested the SDMs with the same dataset.

The conversion of the continuous suitability gradient produced by the SDMs into binary predictions of species distribution requires the choice of a threshold (Elith *et al.*, 2006; Peterson, 2006). The threshold that we chose is derived from the ROC curve. By plotting the sensitivity against 1-specificity for all existing thresholds, the method identifies the value at which the omission and commission errors intersect (Pearce & Ferrier, 2000; Jiménez-Valverde & Lobo, 2007). The models were evaluated using a threshold-dependent method, the True Skilled Statistics (TSS – Allouche *et al.*, 2006; Liu *et al.*, 2011;). The TSS varies from –1 to +1. Negative and near-zero values are no better than random and values near +1 denote the same observed and modelled distributions (Liu *et al.*, 2009). We judged models acceptable only if they had TSS values ≥ 0.5 (Fielding & Bell, 1997). We used the variance equation for TSS proposed by Allouche *et al.* (2006) to calculate the 95% confidence interval for TSS values obtained in this study. We used repeated-measures ANOVAs to compare differences in TSS values of each species using different statistical methods for modelling.

Gap analysis

We based the gap analysis on the presence of a particular set of environmental conditions appropriate to the species occurrence in PAs (Rodrigues *et al.*, 2003). We used the modelling procedure that showed higher TSS values to assess the degree that PAs overlap the distribution of turtle species considered as conservation target.

In Brazil, there are two main categories of PAs: integral protected areas (IPA), which are created for biodiversity preservation and to be free of human interference, and sustainable use areas (SUA) where the sustainable extraction of natural resources is allowed based on management strategies. Each of these types is further divided into various subcategories (SNUC, 2002). In addition, the country has a large percentage of indigenous lands (IL), where indigenous populations have possession and usage rights. We downloaded the official maps of the state and federal PAs from the government website (MMA, 2015) and converted to a resolution of 4 km² for performing the gap analysis.

We ran the analysis considering three kinds of PAs with different levels of protection: (1) IPA only; (2) IPA + SUA; and (3) IPA + SUA + IL. According to Rodrigues *et al.* (2003), the target amount for protecting species should be

related to species range sizes. Small range size species (< 1000 km²) should have 100% of their distributions captured in PAs, and species with large ranges (> 250,000 km²) should have at least 10% of their distributions captured in PAs. Targets for species with intermediate range sizes were based on a logarithmic interpolation between 10% and 100%.

We evaluated the protection targets considering the Brazilian Amazon region, where most turtle species are widely distributed. Thus, we classified species as protected (P) when the target percentage of the distribution size was in fact included within PAs, partial gap (PG) when only a portion of the target percentage was included within PAs; and full gap (FG) when the entire range of the species was outside of the PA network (Rodrigues *et al.*, 2003). For fully aquatic species of turtles, we made a 500 m buffer zone around the Amazonian streams and performed the gap analysis only in this portion of the SDMs.

The annual rates of deforestation in the Brazilian Amazon are concentrated in a region known as ‘arc of deforestation’. To determine whether P, PG and FG species are located in areas that show high anthropic pressure, we overlapped the arc of deforestation with species distribution maps. We obtained the arc of deforestation map from the government website (IBAMA, 2015b).

RESULTS

Species distribution modelling

According to the TSS evaluation method, BIOCLIM produced non-acceptable models for all turtle species (0.0–0.14) (see Appendix S3). GLM generated acceptable models only for *Rhinoclemmys punctularia* and *Podocnemis unifilis* (0.11–0.52). The TSS values for SVM methods ranged from 0.05 to 0.72, producing non-acceptable models for 11 species and acceptable models for five species. Maximum entropy generated acceptable models for 14 species (0.38–0.99) (see Appendix S3). Species that have a more restricted distribution in the Amazon, such as *Acanthochelys macrocephala*, *Mesoclemmys nasuta*, *M. vanderhaegei* and *Rhinemmys rufipes* exhibited the highest TSS values. The confidence interval for the TSS values can be seen in Appendix S3.

Repeated-measures ANOVAs indicated that the best statistical method for modelling in relation to TSS values ($F = 69.052$; $P < 0.05$) was maximum entropy (see Fig. 1).

Gap analysis

Turtle species richness was higher in the sedimentary portion of the Amazon basin, in the Amazon/Solimões River drainage and in the Rio Negro drainage. These basins comprise an important region for freshwater chelonian conservation.

To perform the gap analysis, we used suitability maps produced by the maximum entropy method, because it produced the best TSS values. These suitability maps can be

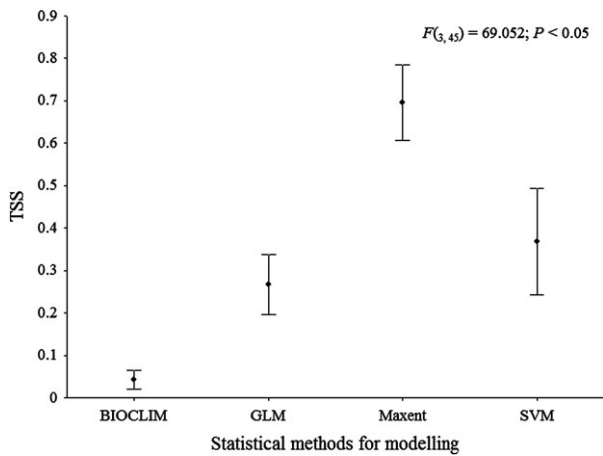


Figure 1 Differences in True Skilled Statistics (TSS) values calculated for turtle species using different statistical methods for modelling.

seen in Appendix S4. We identified only one FG species, *M. nasuta*. The suitable areas for the occurrence of this species were not protected by any category of PA. Other chelonian species were classified as PG species or as fully protected species.

In the highest level of protected area (IPA), only *R. punctularia* and *Kinosternon scorpioides* achieved their protection targets (see Fig. 2a). Thus, IPAs alone do not effectively capture the most suitable areas for turtle occurrence. Under the second level of protected areas (IPA + SUA), we identified 13 species (68.7%) as protected and two species (12.5%) as PG (see Figs 2b and 3b). The PG species occurring in this category of PAs were *M. vanderhaegei* and *A. macrocephala*. These species have the smallest amount of suitable areas in the Amazon, and IPA + SUA were not sufficient to attain conservation targets for them. The species considered fully protected in IPA+SUA scenario had a maximum of 29.2% of their suitable habitat captured in PAs (Fig. 2b, Table 1). Considering all categories of conservation areas (IPA + SUA + IL), *A. macrocephala* and *M. nasuta* were the only species that still did not achieve their conservation targets and were classified as PG species (Fig. 3c). All the other species in this scenario were classified as protected, and they had 27.7–45.1% of their suitable habitat captured by PAs (Table 1).

DISCUSSION

Despite the fact that PAs cover 22.2% the Amazon and IL cover an additional 21.7% (Verissimo *et al.*, 2011), we found some notable gaps in protection of freshwater turtles. The network of integral protection areas is insufficient in capturing the suitable areas for chelonian occurrence. Only *R. punctularia* and *K. scorpioides* are protected by IPAs. These species are semi-aquatic turtles that live in a wide variety of habitats, mostly in small temporary or perennial water bodies in forests. *Kinosternon scorpioides* is a polytypic species

that has a wide distribution, from Mexico to northern Argentina (Rueda-Almonacid *et al.*, 2007; Vogt, 2008). For all other species, we found it was also necessary to consider SUA and IL to reach target protection values, demonstrating the importance of these PA types for effective conservation of freshwater turtles in the Brazilian Amazon.

Our results support the claim that PAs in the Amazon were primarily established to protect terrestrial taxa from overharvesting and deforestation (Peres & Terborgh, 1995; Verissimo *et al.*, 2011). However, such strategies to protect terrestrial species and ecosystems usually do not effectively conserve freshwater ecosystems and their associated fauna (Thieme *et al.*, 2007; Castello *et al.*, 2013). Much of the existing PA network ignores river catchment sites (Wishart & Davies, 2003) and freshwater threats like dams, waterways, oil exploration, pollution (Castello *et al.*, 2013) and flow modification (Abell, 2002; Dudgeon *et al.*, 2006; Davidson *et al.*, 2012; Castello *et al.*, 2013). The mitigation of the impacts of these threats on freshwater ecosystems in Amazon is particularly important because these habitats cover a large area of the basin (Castello *et al.*, 2013) and contribute to the well-being and sustenance of a large number of people (Kvist & Nebel, 2001).

Peres (2005) suggested that megareserves based on biogeographic units defined primarily by the overlap of main river barriers and a vegetation matrix would be adequate to protect Amazon flora and fauna, including aquatic ones. However, we suggest that a catchment-based system for conserving basins would be more appropriate, with identification of areas where terrestrial and freshwater conservation priorities overlap (Castello *et al.*, 2013). Amis *et al.* (2009) noticed that integrating priority areas for conservation of freshwater and terrestrial biodiversity improved management plans in South Africa. Only in particular cases should ecosystems be maintained separately (Thieme *et al.*, 2007). Creating additional PAs in a region where existing PAs already cover a large portion of land is a huge challenge. Thus, a potentially effective strategy for improving protection of freshwater resources would be to prioritize important areas that are also adjacent to existing or proposed PAs, reducing costs (e.g. start-up costs, stakeholder engagement costs) by adding more freshwater biodiversity to existing management efforts (Abell, 2002; Thieme *et al.*, 2007).

Since 1991, most PAs created by the Brazilian government as a policy action for biodiversity protection are sustainable-use reserves (Peres, 2011). Conservation strategies that attempt to reconcile biodiversity conservation and human needs are among the most effective conservation measures (Peres, 2011). However, use of natural resources is often not properly supervised in sustainable-use PAs (Peres & Terborgh, 1995; Peres, 2011). Human pressure induces forest loss, and this impact is one of the major causes of biodiversity loss (Laurance, 1999; Fearnside, 2005). The rural population in Amazon has increased from 6 million in 1960 to 25 million in 2010 (Davidson *et al.*, 2012). Human population densities in Amazonian reserves are frequently larger than in

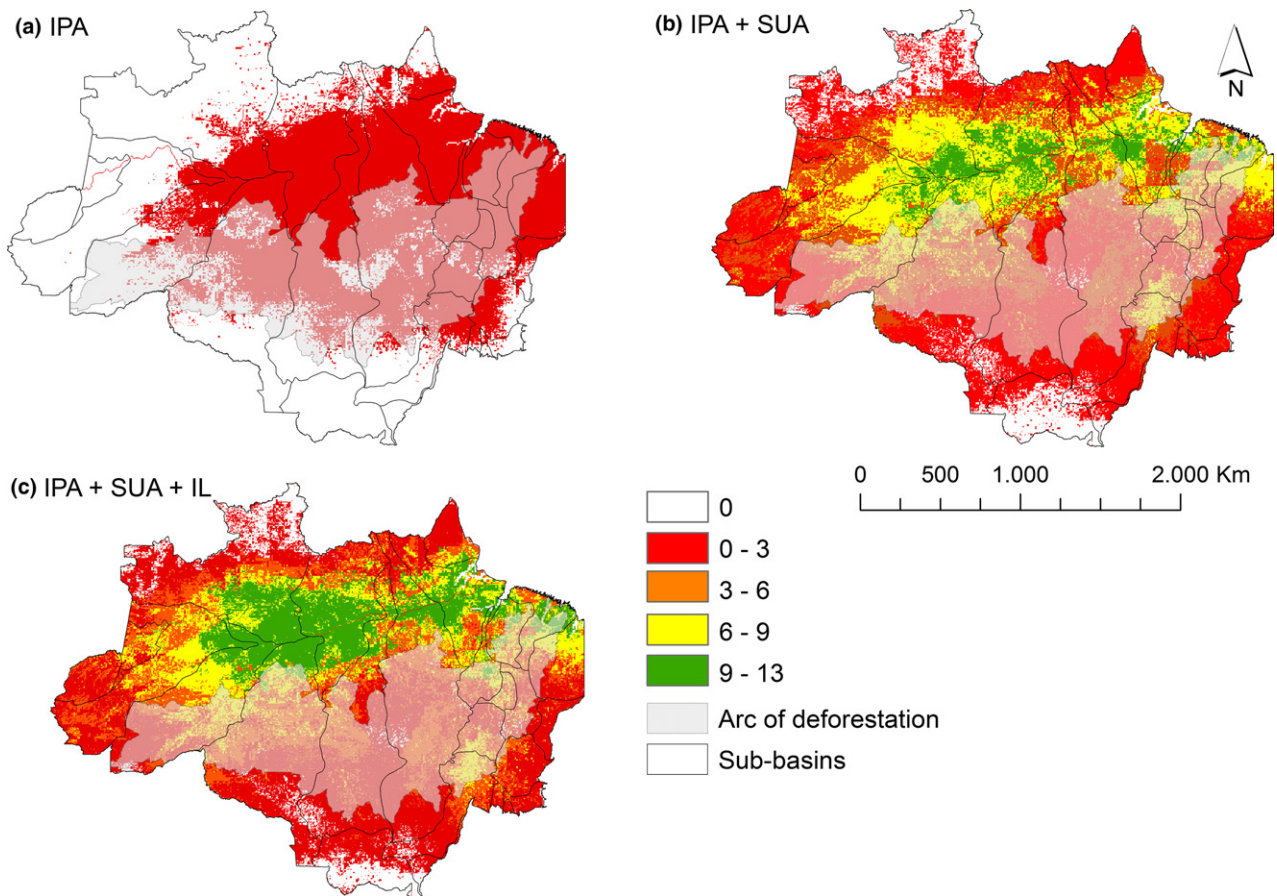


Figure 2 Number of freshwater turtles in Brazilian Amazon fully protected by the reserve networks. The conservation targets are based on the amount of suitable areas generated by maximum entropy method in protected areas. Different levels of protected areas evaluated include the following: (a) integral protection areas (IPA); (b) integral protection areas + sustainable use areas (IPA + SUA); and (c) (integral protection areas + sustainable use areas + indigenous lands (IPA + SUA + IL).

non-PAs (Peres, 2011) and even strictly protected reserves in Brazilian Amazon contain illegal human communities (SNUC, 2002). Since their formal establishment, SUAs have lost 298,500 ha of forest (Veríssimo *et al.*, 2011). Because development in the Amazon is concentrated around waterways, aquatic and semi-aquatic wildlife species are likely heavily impacted (Peres, 2000, 2011).

Conservation success has often been judged by measuring vegetation cover change across large scales (Gaston *et al.*, 2008). The rates of forest loss in Amazon are higher in 'arc of deforestation', a continuous area stretching from the south-west to north-west part of the Amazonian basin (Fearnside, 2005). According to our analysis, suitable areas for several chelonian species occur in this region and are partly captured by the existing PA network, primarily SUAs and ILs. However, turtles may be overharvested even in well forested areas, because hunting is usually unsustainable in an extraction scale (Peres & Lake, 2003). Many populations of game species have been eradicated in extractive reserves (Peres & Palacios, 2007), and chelonians are important in the diet of traditional communities in the Amazon (Kemenes & Pezzuti, 2007; Vogt, 2008; Schneider *et al.*, 2011).

Overcollection of adult females and eggs have been reported as the main threats to the survival of turtle populations, mainly Podocnemididae (Fachín-Terán & Von Mülhen, 2003; Caputo *et al.*, 2005; Fachín-Terán, 2005; Vogt, 2008). One conservative analysis suggested that in the 1980s and 1990s, between 38.79 and 95.11 adults of *P. unifilis* and from 59.15 to 145.02 adults of *P. expansa* were consumed annually by the low-income rural communities in the Brazilian Amazon (Peres, 2000). Hence, sustainable-use reserves may not be sufficient on their own to conserve some freshwater turtles.

According to our analysis, a substantial amount of suitable habitat for species of genus *Podocnemis* is captured in IPAs and SUAs. However, these PAs are not sufficient to capture suitable habitats for species that have restricted distributions in the Brazilian Amazon, such as *M. vanderhaegei*, *M. nasuta* and *A. macrocephala*. *Acanthochelys macrocephala* and *M. nasuta* are not protected in the Amazon, even when we considered all the categories of PAs (IPA + SUA + IL). The distribution of *Acanthochelys macrocephala* in the Amazon is limited to a small part of the south-east region, and the species also occurs in the Brazilian Pantanal, northern Paraguay

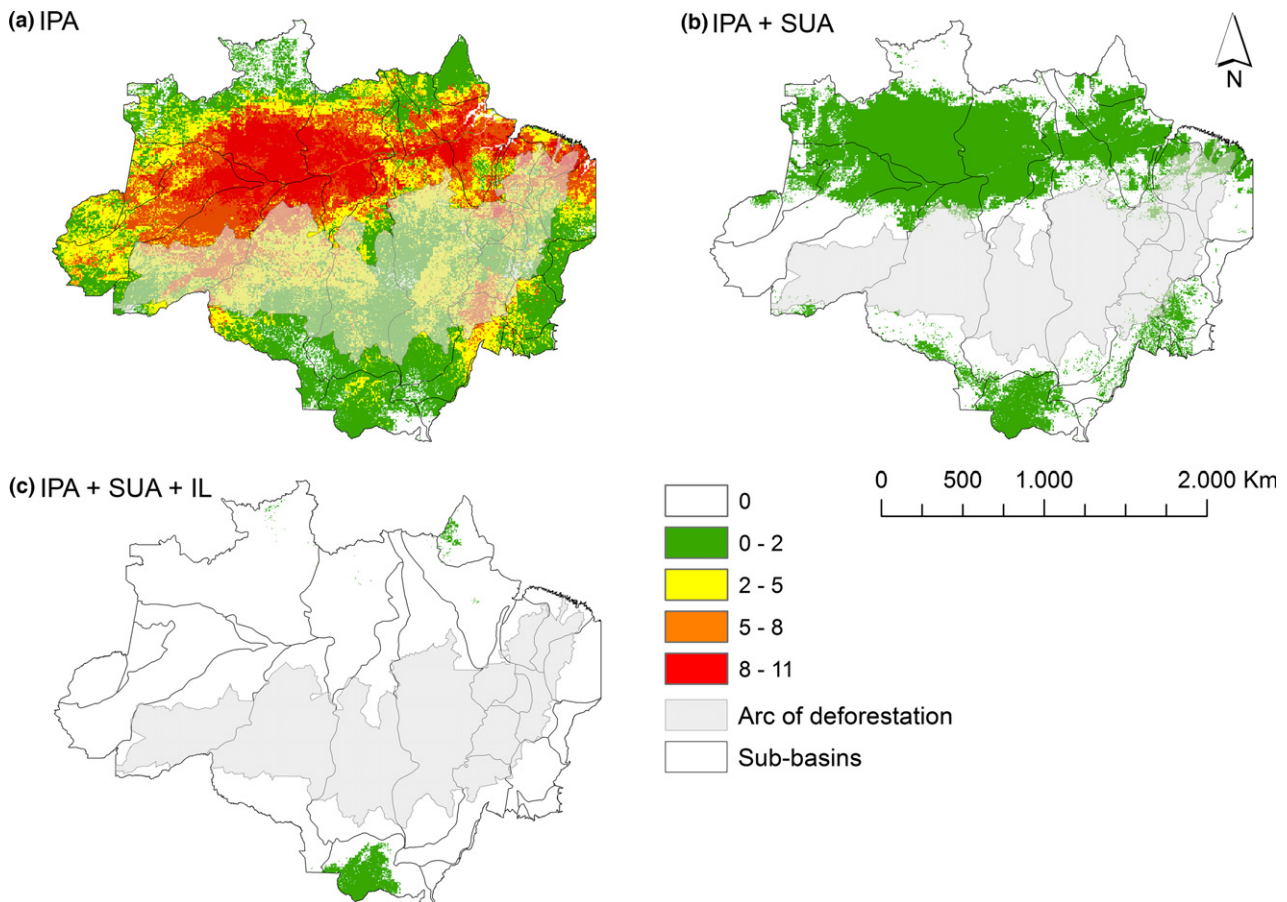


Figure 3 Number of freshwater turtles in Brazilian Amazon that are not protected by the reserve networks (partial gap). The conservation targets are based on the amount of suitable area generated by maximum entropy method in protected areas. Different levels of protected areas evaluated include the following: (a) only integral protection areas (IPA); (b) integral protection areas + sustainable use areas (IPA + SUA); and (c) integral protection areas + sustainable use areas + indigenous lands (IPA + SUA + IL).

and a very small part of Chaco ecoregion in Bolivia, where the effectiveness of PAs could be different (Rhodin *et al.*, 2009). *Mesoclemmys nasuta* is restricted to the Guianas and northernmost Amazon, in the state of Amapá (Bour & Zaher, 2005). Practically, no data concerning the biology and ecology of *M. nasuta* currently exist considering that, until recently, *M. nasuta* was considered conspecific with *M. raniiceps*. Future genetic studies may recombine these allopatric species.

The sedimentary basin in northern Amazon is recognized as an important region in terms of turtle richness, as identified by Buhlmann *et al.* (2009). The area includes priority areas for freshwater turtle conservation. In this region, some of IPAs, such as Reserva Biológica do Rio Uatumã, Reserva Biológica do Rio Trombetas, Reserva Biológica do Abufari and Estação Ecológica de Jutai-Solimões, have already implemented conservation actions for the most impacted species (*P. expansa*, *P. unifilis* and *P. sextuberculata*). Nevertheless, current activities are restricted to environmental education for traditional communities and protection of nesting beaches during the nesting season (Instituto Chico Mendes de

Conservação da Biodiversidade, personal communication; Wildlife Conservation Society Brazil, personal communication). A more local analysis would be an important step for identifying specific sites for protection and specific management actions. Conservation targets should be developed in agreement with local communities and, in most cases, management activities should be carried out by them. According to Peres & Lake (2003), effective community-based conservation requires a capacity-building programme, regulation of immigration into PAs, establishment of sustainable harvest quotas and the creation of intangible zones within reserve boundaries.

In our study, SDMs were useful to predict the geographic range of chelonian species. The distribution of the majority of freshwater turtles in South America is poorly known (Souza, 2004). The predictive capacity of SDMs has been important in addressing urgent conservation problems, especially for rare and unknown species (Pearson *et al.*, 2007; de Siqueira *et al.*, 2009). SDMs have also been critical for rigorous gap analyses and the establishment of conservation priorities (Loiselle *et al.*, 2003; Martinez *et al.*, 2006; Nóbrega &

De Marco Jr., 2011). For particular turtle species, several studies have applied SDMs to help develop conservation policies (Forero-Medina *et al.*, 2012; Ihlow *et al.*, 2012; Millar & Blouin-Demers, 2012). However, the only other study that uses SDMs to generate conservation priorities based on geographic patterns of species richness and vulnerability information for a large group of chelonian species (Trionychidae and Pelomedusidae) was for African freshwater turtles (Bombi *et al.*, 2011).

Comparatively, maximum entropy produced the most reliable SDMs, according to the performance evaluation method we used (TSS). Elith *et al.* (2006) and Pearson *et al.* (2007) suggested that this statistical method is one of the most reliable SDM methods, especially for biased data. However, even using the TSS, which may control for differences in prevalence (Allouche *et al.*, 2006), models for some species, such as *M. raniceps*, *M. gibba* and *Phrynops geoffroanus*, were not acceptable. There are known identification and taxonomic challenges with these species that may contribute to poor model performance. *Phrynops geoffroanus* does not have a clear distribution pattern and is absent only at high southern latitudes (Souza, 2005). The species also seems to be a complex of sibling species (Pritchard & Trebbau, 1984). *Mesoclemmys gibba* has a wide distribution, rather similar to that of *M. raniceps* (Pritchard & Trebbau, 1984; Iverson, 1992b; McCord *et al.*, 2001) and may be misidentified in some occasions (Ferronato *et al.*, 2011). To improve SDMs and conservation planning for these species, we recommend that taxonomic revision efforts be continued for these groups and that new inventory studies be completed.

CONCLUSIONS

Amazonia covers an area of large turtle richness (Buhlmann *et al.*, 2009), composing an important region for their conservation. However, suitable areas for freshwater turtle's occurrence are not protected by the current network of IPA. The insertion of SUA and IL was crucial to consider protected large-range species, but some chelonians may be overharvested in those areas. Facing the current condition, it is necessary to shift the Amazon conservation focus and restructure the PAs in order to contemplate river catchment sites in whole basins. It is necessary to include protection actions that handle the upstream drainage network, the riparian area and, in the case of migratory species, the downstream drainage (Pusey & Arthington, 2003). At this level of PA coverage, not only turtles but all freshwater species would benefit (Dudgeon *et al.*, 2006). The approach would require a new distribution of the PAs and the use of large portions of land as PAs. Thus, a more practical manner to develop a chelonian conservation planning could take into account important areas for turtle richness conservation next to existent PAs and consider features of the traditional cultures in conservation planning in order to attend their needs.

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SUPPORTING INFORMATION

Additional Supporting Information may be found in the online version of this article:

Appendix S1. The 42 environmental variables used for predicting freshwater turtle habitat suitability.

Appendix S2. Summary of the principal components (PCA) used as environmental layers.

Appendix S3. Summary of the evaluation of the species distribution models (SDMs) according True Skilled Statistics (TSS) method.

Appendix S4. Environmentally suitable areas for the occurrence of 16 freshwater turtles in the Amazon.

BIOSKETCHES

The overall aim of this project was to evaluate whether freshwater turtles are protected by the current network of Amazonian conservation units (gap analysis). The lead author of this study is **Camila K. Fagundes**, researcher at Wildlife Conservation Society in Brazil. Her research interests focus on vulnerability of freshwater turtle to land use/land cover changes and species distribution modelling aimed to management practices. **Paulo De Marco Júnior** is currently Associate Professor of the Universidade Federal de Goiás and is permanent advisor in graduate courses of Ecology and Evolution and Environmental Sciences of the cited university. His experience has an emphasis on Theoretical Ecology: community ecology, population ecology, conservation biology and quantitative ecology. **Richard C. Vogt** is a permanent researcher at the National Institute for Amazonian Research (INPA) and advisor in graduate programmes of Tropical Ecology and Freshwater Biology and Inland Fisheries of INPA. His research interests are focused on biology and ecology of the Amazon turtles. He is one of the pioneers in evaluating the effect of incubation temperature on sex determination in turtles.

Author contributions: C.K.F and P.D.M. originally formulated the ideas presented in this study. C.K.F and R.C.V. provided the species data. C.K.F. supplied the environmental data. C.K.F and P.D.M. ran the species distribution models and the gap analysis. C.K.F. wrote the first draft of this manuscript, and P.D.M. and R.C.V. contributed extensively to the preparation of the final version.

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