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# An application of maximum entropy model to evaluate the differential effect of cage aquaculture on the distribution of a native and an endemic fish species in Lake Maninjau, Indonesia 

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#### Abstract

Physical cage aquaculture structure can attract native fish species in marine and freshwater ecosystems. Most studies on the effects of cage farms on native fish distribution have been undertaken in marine environments and outside of Asia as the main freshwater cage aquaculture producing region. Many studies have emphasised connections between native fish distribution and feeding time. Previous research also has shown the necessity to use modelling to monitor this effect to reduce data collection costs. Here we analyse the distribution of an endemic fish species, Rasbora maninjau, and a native fish, Gobiopterus sp., associated with tilapia cage aquaculture occurrence using a Maximum Entropy Model (MaxEnt). We find that the application of the MaxEnt model can produce reliable and accurate information on the impacts of cage aquaculture on the native fish species distribution aligning with the more expensive count data method. Our results also suggest that the species-specific interaction between the native fish and cage farms is mainly arising from an interaction between the ecological behaviour of the native fish with dimensions of the environmental condition such as turbidity. Our study therefore highlights the importance for improved appraisal of the ecology of native fish in the cage aquaculture risk assessment.


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Aquaculture impacts; fish distribution; modelling; freshwater lake; tilapia farm

## 1. Introduction

Growing global demand for fish has led to a rapid increase in freshwater cage aquaculture (FAO 2016; Naylor et al. 2021). In Asia, freshwater bodies in populous fish-eating regions have become centres of intensive cage aquaculture development (Newton et al. 2021; Taskov et al. 2021). Freshwater cage aquaculture can contribute to nutritional security and its function as rural livelihood support has stimulated its growth in some

[^0]countries (Rajee and Mun 2017; Shava and Gunhidzirai 2017; Njiru et al. 2018). However, there are increasing conflicts with other freshwater resource users, particularly where rapid expansion of cage aquaculture has resulted in eutrophication of reservoirs, declines in yields of native biota and increased occurrences of "fish kills" (Abery et al. 2005; Endah et al. 2017; Taskov et al. 2021).

The rapid expansion of freshwater cage aquaculture affects native fish communities in various ways. Firstly, the release of organic matter from feed remains results in increases in primary productivity which can lead to eutrophication, change in water quality (Abdel-Tawwab et al. 2002; Baccarin et al. 2005; Canonico et al. 2005; Chislock et al. 2013; Verdegem 2013) and decrease of benthic fauna richness (Tomassetti et al. 2016). Secondly, native fish communities may be affected by alterations of habitat and water quality and exchanges of parasites and diseases between the native and cultured fish (Barrett et al. 2018; Njiru et al. 2018). Thirdly, native fish populations may be negatively impacted by competition for resources, predation, and hybridization, whether intentionally or accidentally (Barrett et al. 2018).

Recent studies of cage aquaculture in freshwater systems have also investigated the effects of the cage structures on native fish populations (Demétrio et al. 2012; Ramos et al. 2013; Nobile et al. 2018). The physical cage structure may act to aggregate wild fish, offer shelter, and attract wild fish to the continuous food input (Sanchez-Jerez et al. 2011; Nobile et al. 2018). Studies in neotropical reservoirs have suggested that cage structures can increase habitat complexity in freshwater bodies and provide protection from predation for native fish communities (Nobile et al. 2018). However, studies in tropical lakes in other biogeographical regions, for example in Asia as the primary freshwater aquaculture fish producing region, are scarce. Further, previous research identifying species-specific relationships between marine cage farms and native fish species mainly emphasised the influence of feeding time and season (Uglem et al. 2009; Bacher et al. 2012; Ballester-Moltó et al. 2015). Research addressing which factors affect the species-specific relationship between cage farms and native fish distribution in tropical lakes is rare. This study aims at filling these gaps by analysing the effects of tilapia cage farms on a native and an endemic species in Lake Maninjau, Indonesia.

Most research to date aimed at monitoring the cage aquaculture impact on native fish distribution relied on count data collection (Šegvić Bubić et al. 2011; Bacher et al. 2015; Pereira et al. 2019). Collecting count data such as abundance is time consuming and requires specific sampling techniques (Marini et al. 2018; Siddiqui et al. 2018) making it less affordable for lake managers in the study area and in many other areas in Indonesia and Global South Nations where monitoring efforts are limited. Research by Yuniarti et al. (2021) revealed that the lake managers in the study area have voiced concern about limited monitoring resources such as time, funds, and human resources.

In this study, we employ Maximum Entropy Models (MaxEnt) to examine the distribution of two fish species (bada and rinuak) in relation to farm occurrence in Lake Maninjau. MaxEnt requires presence data only and environmental variables which can be collected with relative ease. MaxEnt models have been widely applied in conservation science to predict species distribution and habitat suitability (Sobek-Swant et al. 2012; Moore et al. 2016; Yi et al. 2016; Zhang et al. 2019; Zhang et al. 2018) and has been proven useful when only small sample sizes are available (Elith et al. 2006; Wisz et al. 2008; West et al. 2016). Considering these merits, we test whether the model application can provide a more affordable monitoring approach for cage impacts of tilapia on bada and rinuak, and reflect on the wider applicability of the MaxEnt model approach for other inland water bodies.

## 2. Materials and Methods

### 2.1. Site description

Lake Maninjau (Figure 1) is one of several large volcanic lakes in West Sumatra, Indonesia. The maximum depth in the lake is 165 m . It is 16.6 km from north to south, and 7.5 km at its widest point (Fakhrudin et al. 2002).

Cage aquaculture was introduced to the lake in 1992 with the installation of 16 cage units which produced 96 tons of fish in three months (Syandri and Azrita 2013). The currently farmed fish species on Lake Maninjau is Nile tilapia (Oreochromis niloticus). The number of cage units present in Lake Maninjau has increased rapidly, with 4,000 further units established in 2003, totalling 23,566 in 2016 (Syandri et al. 2014; Agam Regency Fisheries Department 2017). (The increasing presence of cage aquaculture in the lake has caused concern that aquaculture development has exceeded the lake's carrying capacity and accelerated eutrophication (Said et al. 2020).

Long term satellite data showed that cage aquaculture proliferation has caused significant decline of the lake's water transparency (Setiawan et al. 2019). Further, annual yet patchy data of the lake's water quality revealed that the expansion of cage aquaculture operation is correlated with an decrease of oxic layer, an increase of total phosphorus concentration, and an elevation of trophic state index (Sulastri et al. 2015).

### 2.2. The fish species

The main fish catch of the lake consists of two benthopelagic fish species: Rasbora maninjau (Lumbantobing 2014) (local name: bada) and Gobiopterus sp. (local name: rinuak). Bada is an important endemic fisheries species and a local culinary icon (Dina et al. 2019). Meanwhile, rinuak, is an important native fish for local communities around Lake Maninjau.

Studies on these two fish species, particularly on rinuak, are scarce and limited. Experts have not yet agreed on the classification and species name of rinuak. The species name for rinuak had not been assigned at the time of writing. A study by Roesma et al. (2020) using, NCBI database (GenBank) described rinuak as a close relative to Gobiopterus, but did not specifically classify its genus. Further, research by Larashati (2019, unpublished data) using the BOLD database showed that it belongs to Gobiopterus. Moreover, its biological and ecological characteristics remain understudied (Kottelat et al. 1993). More importantly, its catch rates remain largely unknown. To the best of our knowledge, rinuak was known to be susceptible to tubo belerang - a local name for the turnover of water


Figure 1. Study site (Image is processed in QGIS 3.2.1 based on map provided by, retrieved on 2 March 2021).
and toxic material (e.g. $\mathrm{H}_{2} \mathrm{~S}$ ) from hypolimnion to epilimnion layer adding more severe impacts to anoxia condition due to eutrophication (Yuniarti et al. 2021b). It is also susceptible to the concentration of ammonia in the water (Yoga and Samir 2021).

On the contrary, bada shows higher survival capability during tubo belerang due to its agility and its strong connection with riverine waters, because they need to migrate when they are spawning (Hartoto and Mulyana, 1996 cited in Dina et al. 2019; Dina et al. 2019). Bada's diet, which primarily consists of zooplankton, aquatic insects, and small portions of phytoplankton, was known to overlap with the food of wild tilapia (Yuniarti and Sulastri 2010; Dina et al. 2019). It is reported that in recent years there has been a declining trend of bada production due to marble goby predation and overfishing (Dina et al. 2019; Linggi et al. 2020).

### 2.3. Selection of environmental variables

To select the environmental variables (EVs) for the models, we adopted the approach presented by Yiwen et al. (2016). Owing to the lack of ecological knowledge about bada and rinuak, we selected several EVs based on general fish ecological knowledge. The EVs that were selected a priori were cage presence, turbidity, presence of natural canopy, dissolved oxygen (DO), pH , conductivity, and water surface temperature (see Rosette et al. 2020)

### 2.4. Data collection and availability

Prior to conducting the habitat survey, a focus group discussion with local fishers was conducted to ensure that we also include the natural habitat of the fish in the survey. We also engaged one of the fishers to guide us during the survey.

A survey of the presence and absence of the fish and measurement of environmental variables was conducted in April 2019. An underwater camera (GoPro Hero 7) was used to survey fish presence. The survey was conducted from 9:00 am to $5: 00 \mathrm{pm}$ for four days for each species representing feeding and non-feeding time. Fish presence was recorded from the water surface $(0 \mathrm{~cm})$ and in different depths depending on habitat type. We selected four habitat types based on information gathered from interviews from eleven fishers. Surveys were conducted in: (1) leafy canopy covered lake margins (depth of underwater video: $0 \mathrm{~cm}, 30 \mathrm{~cm}$, and the lake's bed); (2) the cage farm area (depth of the underwater video: $0,30,60,90$ and 130 cm - the average secchi depth in cage farm area at the time of sampling); (3) open waters 1 km from lake's edge (depth of the underwater video: $0,30,60,90$, and 175 cm - the average secchi depth in open water area, no canopies, no cages); (4) shallow water with no canopy ( 0 and 23 cm the lake's bed). The water depth was measured using a secchi disk stick. However, we did not record any evidence fish presence below 50 cm . In total, we surveyed 36 sites for presence and absence of rinuak (Figure 2(a) and Table A), 23 sites for bada's presence (Figure 2(b) and Table B), and 33 sites for environmental variables (EVs) (Figure 2(c)).

We conducted a second field survey in March 2020 to collect data on presence and absence (Go Pro sampling) for independent cross validation referring to the work of West et al. (2016). The presence and absence data obtained from this second survey was used to validate models by providing ground data of the fish distribution. The survey was conducted randomly at 40 sites (Tables C and D). The random sampling sites were generated by using a research tool in QGIS by determining 1,000 metres as the minimum distance between points.

To verify that MaxEnt can be useful to assess the cage attraction to the fish compared to the usual method using count data, we conducted sampling of fish abundance in the


Figure 2. Sampling location (a) Rinuak, (b) Bada, (c) Environmental variables.


Figure 3. Relative abundance sampling locations (a) Rinuak, (b) Bada.
same month. The sampling was done following the work of Brandão et al. (2013). We used the same habitat criteria as for the presence/absence sampling to determine the abundance sampling sites (native/control habitat, the farm area, and open waters). Rinuak were caught using a scoop net with three repetitions at each site (Figure 3(a)). It was sampled in the morning between 6:00 am to 11:00 am to coincide with first light and during the period of maximum fish activity as informed by the local fishers.

Abundance data sampling of bada (Figure 3(b)) was conducted by experienced fishers using gill nets (size $100 \times 3 \mathrm{~m}$, mesh size 19, 25, and 38 mm ) in April 2019. Multiple mesh size gill nets were recommended to sample fish from freshwater and estuarine areas (Gray et al. 2005). The nets were set from 2:00 am to 5:00 am following the guidance of the fishers and is the optimum period to catch this species, and not the farms' feeding time. The sampling was repeated for three consecutive nights. The data is provided in Tables C and D. All data are available in https://github.com/ivanayuniarti02/Maxent.

### 2.5. Data analysis

Before building the models, to test if the EVs data are spatially correlated, we calculated local Moran's I Index (I) using autoregressive STATA 15 (Anselin 1995; Kondo 2021). Local Moran's I index shows the indication of spatially autocorrelated environmental variable data, which may be caused by selecting sampling points that are too close to each other. In other words, it is used to test if the data is affected by the observation of a nearby sampling site (Rousset and Ferdy 2014). The I indices and their p-values were calculated and presented for DO, pH , turbidity, conductivity, and temperature.

To test the effects of the spatially autocorrelated EVs data on the model performance, we built two sets of models for each species. The first set (referred as total variables/ TVM) included all EVs. The second set (partial variables/PVM) was constructed by removing those EVs which are indicated to be spatially autocorrelated.

In the model building process, we ran the models using presence only data for both species, because we do not have absence data for bada. MaxEnt automatically generates predicted absence data using the environmental predictors' information to predict the absence of the species referred as pseudo-absence (Phillips and Dudi'k 2008). We run the model in default settings/auto features mode (linear, quadratic, product, hinge) to obtain the best fitting model following the work of Phillips and Dud'k (2008), Jose et al. (2020), Yiwen et al. (2016).

Specifically, we set the regularization multiplier (RM) to avoid overfitting - a modelling error which happens when a function of the model relates too closely to a particular set of data looking the model too closely. We specified the RM at $0.5,1.0$, 1.5, 2.0, 4.0 for both model sets (Jose et al. 2020). RM is a tool in MaxEnt which can be used to regulate how focused or closely-fitted the output distribution is (Waszkowiak et al. 2002). Further, to obtain a stable model, we used a 5 -fold cross-validation (CV) approach for both species in each model set (TVM or PVM). CV is a method to select the optimal model when the data is limited (De Bin et al. 2016)

We categorized the model settings as type 1 (CV: 5; RM:0.5), type 2 ( $5, \mathrm{RM}: 1$ ), Type 3 (5; RM:1.5); type 4 ( $5 ; \mathrm{RM}: 2$ ), and type 5 ( $5 ; \mathrm{RM}: 4$ ). Models were run with forty replications. Ten thousand random background data/grid points were selected to generate the model. In total, we constructed a total of 25 models for each species and each model set ( 5 models for each RM).

To compare models' performance and to select the best fitting model, we used the Area Under Curve presence only $\left(\mathrm{AUC}_{\mathrm{PO}}\right)$ indicator. The $\mathrm{AUC}_{\mathrm{PO}}$ is produced in MaxEnt using presence only and pseudo-absence data (Yackulic et al. 2013).

To validate the models using independent field data (obtained from the second presence data sampling), we calculated several indices, including sensitivity, specificity, and the True Skill Statistic (TSS) (Allouche et al. 2006; West et al. 2016) for the best fitting models (setting type 3 for both species). A cut-off point of 0.5 was used in this step to obtain predicted occurrence (probability $\geq 0.5$ means presence, and otherwise).

To find the most influential environmental parameters, we used a stepwise removal method (Whittingham et al. 2006; Yiwen et al. 2016). The stepwise removal step was done on the best fitting models. We relied on the Jackknife analysis to generate indicators of the environmental predictors' importance. The process was repeated until two variables were obtained, because two variables are the minimum requirement for input data resulting in four level models for each species. Again, we used $\mathrm{AUC}_{\mathrm{PO}}$ to evaluate the performance of the model performance (Phillips et al. 2016; Yang and Berdine 2017)

In the end, to test if the results of MaxEnt aligns with the results of the count data method, we used a maximum likelihood mixed effect model (ML) drawing on the relative abundance data - numbers of individual per species per square metre of area. We used STATA 15 to perform the test. ML was selected because the model can analyse the impact of repeated measurement (pseudo-replication). We tested the models against the null model. The Akaike Information Criteria (AIC) and Likelihood Ratio Test (LRT) were used to compare the models.

## 3. Results

### 3.1. Spatial autocorrelation of environmental variables

The local Moran's I indices (I) and the expected indices (E(I)) are generated to calculate the p -value (Table 1). The I values show that DO and pH are spatially autocorrelated, which means that the measurement of these parameters is influenced by the adjacent sampling sites. The p-values suggest that we can reject the null hypothesis that there is no spatial autocorrelation for both EVs. Further, the I values for both EVs are above zero indicating positive SAC, which means that the observation of DO and pH in one sampling point is affected by the results of observation in the nearby point. Meanwhile, SACs for the other three EVs are not detected. Considering the results, we built the models with and without these parameters to test whether removing the EVs with SAC would affect the models' performance.

### 3.2. Model selection

The estimated test statistics of Area Under ROC Curve (AUC) are quite similar for both model sets (Partial Variable Models/PVM and Total Variable Models/TVM). The values above 0.9 mean the models have good predicting capability. Therefore, that removing the spatially autocorrelated EVs does not dramatically change the model performance (Table 2).

The selected final models were partial variable models (PVM) to ensure that the explanatory variables were not spatially autocorrelated. Based on the optimum AUC values and the resulting maps, we selected PVM with model setting type 2 ( 5 replications with regularization multiplier 1) for rinuak. PVM with model setting type 1 (5 replications with regularization multiplier 0.5) is chosen for bada.

Drawing on the field validation using independent test data true statistic skill values (TSS) are estimated to be above zero indicating that the models performed better than random prediction for both species (Table 3) (Allouche et al. 2006; West et al. 2016).

The calculated TSS values reveal that the model is able to predict the actual occurrence of the fish as shown in Tables E and F. Although some predictions of the presence deviate from actual occurrence, most of the predicted absence aligns with the observed data.

Table 1. The local Moran's I index of the environmental variables presenting spatial autocorrelation in DO and pH .

| Environmental |  |  |  |  |
| :--- | :---: | :---: | :---: | :---: |
| variable | I | $\mathrm{I})$ | Z | p -value |
| $\mathrm{DO}^{\mathrm{a}}$ | 0.534 | -0.029 | 3.899 | $p<0.05$ |
| $\mathrm{pH}^{\mathrm{a}}$ | 0.581 | -0.029 | 4.224 | $p<0.05$ |
| Turbidity | 0.037 | -0.029 | 0.519 | 0.302 |
| Conductivity | 0.009 | -0.029 | 0.274 | 0.392 |
| Temperature | 0.040 | -0.029 | 0.527 | 0.311 |

Note: $\mathrm{H}_{0}$ : spatial randomness
${ }^{\text {a }}$ statistically significant, indicating occurrence of spatial autocorrelation

Table 2. AUC values of the constructed models to choose the fittest models.

| Species | Model settings | Total variable models (TVM) ${ }^{\text {c }}$ | Partial variable models (PVM) ${ }^{\text {d }}$ |
| :---: | :---: | :---: | :---: |
|  |  | Test $\mathrm{AUC}_{\text {PO }}{ }^{\text {a }}$ | Test AUC ${ }_{\text {PO }}{ }^{\text {a }}$ |
| Rinuak | Type 1 (Cross validation: 5, Regularization coefficient: 0.5 ) | 0.928 | 0.928 |
|  | Type 2 (Cross validation: 5, Regularization coefficient: 1) ${ }^{\text {b }}$ | 0.943 | 0.930 |
|  | Type 3 (Cross validation: 5, Regularization coefficient: 1.5) | 0.940 | 0.933 |
|  | Type 4 (Cross validation: 5, Regularization coefficient: 2) | 0.937 | 0.933 |
|  | Type 5 (Cross validation: 5, Regularization coefficient: 4) | 0.920 | 0.927 |
| Bada | Type 1 (Cross validation: 5, Regularization coefficient: 0.5$)^{\text {b }}$ | 0.952 | 0.977 |
|  | Type 2 (Cross validation: 5, Regularization coefficient: 1) | 0.948 | 0.947 |
|  | Type 3 (Cross validation: 5, Regularization coefficient: 1.5) | 0.952 | 0.943 |
|  | Type 4 (Cross validation: 5, Regularization coefficient: 2) | 0.951 | 0.936 |
|  | Type 5 (Cross validation: 5, Regularization coefficient: 4) | 0.935 | 0.919 |

Note: ${ }^{\mathrm{a}} \mathrm{AUC}_{\text {PO }}$ (AUC Presence Only) as proposed by (Yackulic et al. 2013) to indicate that the AUC value is produced as a part of MaxEnt model building using presence only and pseudoabsence data.
${ }^{\text {b }}$ The selected model settings
'Total variable models (TVM): the models built with all included environmental variables. dPartial variable models (PVM): the models built with removing spatially correlated environmental variables ( DO and pH ).

Table 3. Results of independent data field validation showing the good prediction ability of the models.

| Species | Sensitivity | Specificity | TSS |
| :--- | :---: | :---: | :---: |
| Rinuak | 0.60 | 0.80 | 0.40 |
| Bada | 0.70 | 0.71 | 0.41 |

### 3.3. Species distribution maps

The maps (Figure $4(\mathrm{a}, \mathrm{b})$ ) present the predicted pattern of occurrence of both species in response of the environmental variables (cage, temperature, turbidity, conductivity, and canopies). Overall, predicted presence decreases with the distance from the lake's edge especially for rinuak. This species was mapped to be present mostly on the lake's edge compared to bada, and almost overlapped with the cage farms (Figure 4(a) vs. Figure 4(c)). Meanwhile, although there was overlap in some spots, bada was also predicted to occur in the more open waters (Figure 4(b,c)).

### 3.4. The important environmental variables affecting fish distribution

The results of the stepwise analysis shown in Table 4 are used to identify the principal environmental parameters influencing the occurrence of both species. Overall, temperature is the most influential variable for both rinuak and bada and turbidity ranks as the second most important variable for bada, and the third for rinuak. (Table 5). Meanwhile, cage presence, the second most influencing variable for rinuak distribution, is not closely associated to bada's presence.

### 3.5. Relative abundance in three habitat types

We could only find rinuak in leafy canopied and cage areas as the fish are absent in open water areas. Thus, we only use those two habitat types in the ML model for rinuak (Figure 5(a)). Meanwhile, the habitat types tested for bada are leafy canopied habitat, cage farms, and open waters as the fish were present in those three habitat types (Figure 5(b)).

The results of the mixed effects model (Table 5) show that only habitat types have significant influence on rinuak's relative abundance (Likelihood Ratio Test against null model/ LRT $^{1}<0.05$ and LRT models with habitat and sampling location as variables and models


Figure 4. Maps of predicted species distribution in correlation with cage presence, red, yellow, and green colours show the predicted presence of the species (a) Rinuak, (b) Bada, (c) Cage farm.

Table 4. Results of stepwise analysis to obtain the most important environmental variables.

| Species | Ranking of included variables | Test AUC ${ }_{\text {PO }}$ | Remark |
| :--- | :--- | :---: | :--- |
| Rinuak | Te, Ca, Tur, Con, Can | 0.930 | Excellent determination |
|  | Te, Ca, Tur, Con | 0.900 | Excellent determination |
|  | Te, Ca, Tur | 0.921 | Excellent determination |
|  | Te, Ca | $0.946^{\text {a }}$ | Excellent determination |
| Bada | Te, Tur, Con, Can, Ca | 0.949 | Excellent determination |
|  | Te, Tur, Con, Can | 0.947 | Excellent determination |
|  | Te, Tur, Con | $0.957^{a}$ | Excellent determination |
|  | Te,Tur | 0.942 | Excellent determination |
|  |  |  |  |

Note: Te: temperature; Ca: cages; Tur: turbidity; Con: conductivity; and Can: natural canopy ${ }^{\text {a The }}$ best models

Table 5. Results of maximum likelihood mixed effect model (ML) for rinuak and bada.

| Species | Response Variable | Mixed effects model | df | AIC | LL | LRT ${ }^{\text {b }}$ | LRT ${ }^{\text {c }}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Rinuak | Fish's relative abundance | Habitat + Sampling Locations | 4 | 363.26 | -177.63 | $p=0.04$ | $p=1.00$ |
|  |  | Habitat ${ }^{\text {a }}$ | 3 | 361.26 | -177.63 | $p=0.01$ |  |
|  |  | Null model | 2 | 365.58 | -180.79 | - |  |
| Bada | Fish's relative abundance | Habitat + Sampling Locations | 4 | 52.81 | -22.41 | $p=0.17$ | $p=0.10$ |
|  |  | Habitat | 3 | 53.48 | -23.74 | $p=0.34$ |  |
|  |  | Null model ${ }^{\text {a }}$ | 2 | 52.39 | -24.20 | - |  |

[^1]

Figure 5. Relative abundance: (a) Rinuak, (b) Bada.
with habitat only variable/LRT $\left.{ }^{2}>0.05\right)$. The results for bada indicate no statistically significant effect of habitat types and sampling locations on its relative abundance (LRT ${ }^{1}<0.05$ ).

## 4. Discussion

Water temperature is the most influential variable affecting the distribution of both species. Few studies have addressed the impacts of warming temperature to fish distribution in tropical lakes (Cohen et al. 2016; Barbarossa et al. 2021). In general, rising temperature affects tropical freshwater fish distribution by decreasing their fitness due to increased stress physiology and required energy to reproduce and to survive, which forces the fish to be more actively find a suitable environment (Nivelle et al. 2019; Alfonso et al. 2020). The ability to find the suitable environment is constrained by limited habitat availability (Alfonso et al. 2020; Nivelle et al. 2019). Further, the warming temperature affects habitat availability by reducing water levels, jeopardising habitat connectivity, and disrupting riverine habitat (Ficke et al. 2007; Miranda et al. 2020; Barbarossa et al. 2021). The effect of rising temperature on habitat connectivity will especially affect bada, which migrates to riverine habitat for spawning. Additionally, the warming temperature will cause water retention changes, exacerbate eutrophication leading to anoxia condition, and increase the toxicity of pollutants (Ficke et al. 2007; Missaghi et al. 2017; Miranda et al. 2020). Thus, the impacts of increasing temperature on eutrophication will particularly affect rinuak, whose survival has been threatened by the tubo belerang event (Yuniarti et al. 2021a).

Moreover, our data corroborate that turbidity affects both species, and especially bada, which aligns with the study of Dina et al. (2019), which mentioned that bada prefers clean water. Turbidity may reduce bada's visual ability to spot zooplankton as their main diet (cf. Sulastri et al. 2010; Yuniarti and Sulastri 2010). Zooplankton can easily camouflage in the turbid water impeding bada's foraging behaviour, as indicated by Lunt and Smee (2015). On the contrary, turbid water around the cages could be beneficial for rinuak to avoid predators due to its tiny translucent body. Turbid water and cage structures, which increase habitat complexity, help rinuak to reduce its predation risk by decreasing encounter rates and foraging ability of its predators as explained by Ajemian et al. (2015) and McElroy et al. (2018). Therefore, it is not surprising that small fish like rinuak occupy complex littoral habitat or artificial structures and avoid open water habitat (Ajemian et al. 2015; Merz et al. 2021).

Turbidity is affected by the more frequent extreme weather predicted in Indonesia due to the warming temperature (Measey 2010; Supari et al. 2017). The anticipated longer wet season and increased rainfall will increase sediment run-off will add more debris avalanches to the lake, which is naturally high given the occurrence of earthquake and heavy rain in the study area (Wils et al. 2021). Turbidity was also predicted to be boosted by increasing human activities in the lake's watershed area (Antomi et al. 2016). In addition to that, a strong connection between Lake Maninjau's increased turbidity with the cage activities using long term satellite data has been identified (Setiawan et al. 2019).

Considering that natural and anthropogenic factors influence environmental variables and affect fish distribution, we underline the importance of continuing monitoring to help mitigation planning, as suggested by Callier et al. (2018) and Miranda et al. (2020). We suggest using MaxEnt to overcome monitoring constrains because it has shown good prediction ability of the fish distribution in our case study, making it useful to guide ecosystem managers to focus scientific effort as suggested by West et al. (2016) and Pearson et al. (2007). The MaxEnt models, which can be built based on species presence and environmental variables data, help reduce sampling cost. Using MaxEnt also requires lower specific skills to collect the data than common count data collection. Moreover, MaxEnt models can also be developed to forecast the dynamics of fish distribution related to changing environmental variables, such as changing temperature driven by future climate scenarios (Qin et al. 2017; Borzée et al. 2019; Hadgu et al. 2019).

Further, the Jackknife analysis feature in MaxEnt is beneficial to illustrate the significance of various environmental variables on fish distribution, including cage farming, one of the main anthropogenic activities in the study area. The results of the Jackknife analysis indicate that rinuak presence is highly related to the presence of the cage farms. However, the same does not apply for bada's distribution. We argue that the cage farms offer additional habitat to rinuak because the cage farms are not built under leafy canopies and expand up to 1 km from the lake's edge. The cage farms' attraction to rinuak is related to the habitat services provisioning, as suggested by TEEB (2010). The services offered by the farms are shelter and food source (Sanchez-Jerez et al. 2011; Goodbrand et al. 2013; Ramos et al. 2013; Uglem et al. 2014; Nobile et al. 2018). The farm's sheltering function is primarily associated with the provisioning of artificial physical structures acting like fish aggregating devices or artificial reefs (Dempster et al. 2011; Sanchez-Jerez et al. 2011; Uglem et al. 2014; Barrett et al. 2018; Callier et al. 2018).

The degree of association between wild fish distribution and farm cages is varied and species-specific (Sanchez-Jerez et al. 2011; Arechavala-Lopez et al. 2015; Ballester-Moltó et al. 2015). Our results support this finding by providing evidence that rinuak presence is strongly associated with cage farms, but not bada. As aforementioned, rinuak is likely to benefit from the complex structure of cage farms, waste feed, and turbid water, while bada does not rely on these aspects. Therefore, we suggest that the species-specific relationship between the cage farms and wild fish spatial distribution depends on the interaction between the ecology of the species and the environmental variables such as turbidity. This finding complements previous studies of cage farms' attraction to wild fish, which highlighted that the wild fish-cage farm relationship was affected by feeding time, fish' life stage, and seasonal event (e.g. reproductive cycle) (Sudirman et al. 2009; Uglem et al. 2009; Sanchez-Jerez et al. 2011; Šegvić Bubić et al. 2011; Bacher et al. 2012, 2015; Ballester-Moltó et al. 2015).

Our Data has limitations that should be addressed in future research. First, collecting data at various points in the year is advisable considering that cage aquaculture operates all year round in the lake. Second, future research on similar topics should include the effects of nutrient concentration in the lake's water, including total nitrogen, total phosphorus, and ammonia nitrogen, because these parameters are more sensitive to
aquaculture activities. They may have impacts on native fish distribution through the occurrence of harmful algae bloom and DO depletion.

## 5. Conclusion

A species distribution model such as MaxEnt is shown to provide reliable information of the impacts of cage aquaculture on the distribution of native species compared to the more time consuming and expensive method using count data. Thus, we recommend using MaxEnt to help monitoring efforts in areas with limited monitoring capacities. The model can also be extended to advice the suitable area to move the cage aquaculture related to the future native fish distribution considering the warming temperature and increased turbidity. Further, we also emphasise the importance of an improved understanding of the ecology of native fish, including their interaction with café farms, for conducting environmental risk assessments of cage aquaculture.

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The authors declare that there is no conflict of interests.

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## Supplementary materials

Table A. Presence-absence data for Rinuak.

| Long | Lat | Presence $=1 /$ Absence $=0$ |
| :--- | :--- | :---: |
| 100.1575 | -0.27947 | 1 |
| 100.1588 | -0.27964 | 1 |
| 100.1603 | -0.65267 | 1 |
| 100.161 | -0.27747 | 1 |
| 100.1629 | -0.27333 | 1 |
| 100.1613 | -0.28631 | 1 |
| 100.1678 | -0.27219 | 1 |
| 100.1633 | -0.29089 | 1 |
| 100.1706 | -0.26308 | 1 |
| 100.1863 | -0.25492 | 1 |
| 100.1896 | -0.25917 | 1 |
| 100.209 | -0.26281 | 1 |
| 100.2257 | -0.29392 | 1 |
| 100.2237 | -0.30147 | 1 |
| 100.2251 | -0.31125 | 1 |
| 100.2255 | -0.31125 | 1 |
| 100.2241 | -0.31956 | 1 |
| 100.2241 | -0.32603 | 1 |
| 100.2178 | -0.33319 | 1 |
| 100.1596 | -0.28472 | 1 |
| 100.1558 | -0.28481 | 1 |
| 100.1643 | -0.29983 | 1 |
| 100.1659 | -0.30856 | 1 |
| 100.1665 | -0.32233 | 1 |
| 100.1652 | -0.33172 | 1 |
| 100.1652 | -0.3445 | 1 |
| 100.1797 | -0.35136 | 1 |
| 100.1834 | -0.35569 | 1 |
| 100.1746 | -0.35967 | 1 |
| 100.1694 | -0.37956 | 1 |
| 100.1659 | -0.38611 | 1 |
| 100.1862 | -0.4006 | 0 |
| 100.1912 | -0.33169 | 0 |
| 100.1878 | -0.3445 | 0 |
| 100.1935 | -0.35342 | 0.37 |
|  |  | 0 |
|  | -0.1905 |  |

Table B. Presence-absence data for Bada.

| Long | Lat | Presence $=1 ;$ Absence $=0$ |
| :--- | :---: | :---: |
| 100.1655 | -0.32767 | 1 |
| 100.1538 | -0.29108 | 1 |
| 100.1819 | -0.25942 | 1 |
| 100.1911 | -0.25458 | 1 |
| 100.2111 | -0.26339 | 1 |
| 100.1665 | -0.32233 | 1 |
| 100.1659 | -0.30856 | 1 |
| 100.1797 | -0.35136 | 1 |
| 100.209 | -0.26281 | 1 |
| 100.2255 | -0.31125 | 1 |
| 100.1706 | -0.26308 | 1 |
| 100.1575 | -0.27947 | 1 |
| 100.1652 | -0.33172 | 1 |
| 100.1628 | -0.29753 | 1 |
| 100.1638 | -0.30436 | 1 |
| 100.1638 | -0.31325 | 1 |
| 100.1713 | -0.29028 | 1 |
| 100.1755 | -0.30594 | 1 |
| 100.1686 | -0.30986 | 1 |
| 100.1734 | -0.33364 | 1 |
| 100.1633 | -0.32247 | 1 |
| 100.2257 | -0.29089 | 1 |
| 100.2255 | -0.29392 | 1 |
| 100.1613 | -0.31125 | 1 |

Table C. Abundance data of Rinuak.

| Location | Repetition | Relative abundance (fish/m2) | Type of habitat |
| :---: | :---: | :---: | :---: |
| S 0.2847 E 100.1596 | 1 | 176.30 | Cage area |
|  | 2 | 40.60 | Cage area |
|  | 3 | 78.50 | Cage area |
| S 0.2998 E 100.1643 | 1 | 456.00 | Cage area |
|  | 2 | 69.20 | Cage area |
|  | 3 | 64.60 | Cage area |
| S 0.3445 E 100.1652 | 1 | 61.80 | Cage area |
|  | 2 | 60.00 | Cage area |
|  | 3 | 86.80 | Cage area |
| S 0.3861 E 100.1659 | 1 | 46.20 | Cage area |
|  | 2 | 21.20 | Cage area |
|  | 3 | 27.70 | Cage area |
| S 0.3858 E 100.1842 | 1 | 97.80 | Cage area |
|  | 2 | 325.80 | Cage area |
|  | 3 | 90.50 | Cage area |
| S 0.2981 E 100.1558 | 1 | 97.04 | Littoral zone |
|  | 2 | 28.40 | Littoral zone |
|  | 3 | 56.80 | Littoral zone |
| S 0.3086 E 100.1659 | 1 | 39.23 | Littoral zone |
|  | 2 | 96.92 | Littoral zone |
|  | 3 | 56.92 | Littoral zone |
| S 0.4984 E 100.1652 | 1 | 69.23 | Littoral zone |
|  | 2 | 21.54 | Littoral zone |
|  | 3 | 43.08 | Littoral zone |
| S 0.3317 E 100.1652 | 1 | 150.77 | Littoral zone |
|  | 2 | 87.69 | Littoral zone |
|  | 3 | 212.31 | Littoral zone |
| S 0.3534 E 100.1839 | 1 | 16.92 | Littoral zone |
|  | 2 | 27.69 | Littoral zone |
|  | 3 | 55.38 | Littoral zone |
| S 0.2859 E 100.1880 | 1 | 0.00 | Open waters |
|  | 2 | 0.00 | Open waters |
|  | 3 | 0.00 | Open waters |
| S 100.1917 E 0.3329 | 1 | 0.00 | Open waters |
|  | 2 | 0.00 | Open waters |
|  | 3 | 0.00 | Open waters |
| S 0.3348 E100.1889 | 1 | 0.00 | Open waters |
|  | 2 | 0.00 | Open waters |
|  | 3 | 0.00 | Open waters |
| S 0.3564 E 1001935 | 1 | 0.00 | Open waters |
|  | 2 | 0.00 | Open waters |
|  | 3 | 0.00 | Open waters |
| S 0.3716 E 100.1917 | 1 | 0.00 | Open waters |
|  | 2 | 0.00 | Open waters |
|  | 3 | 0.00 | Open waters |

Table D. Relative abundance data of Bada.

|  |  |  | Relative abundance (fish/m2) |  |  |
| :--- | :---: | :---: | :---: | :---: | :---: |
| No | Location | Repetition | Cage culture area | Littoral zone | Open waters |
| 1 | S 00 15.116' E 100 11.759' | $1(26-02-19)$ | 0.08 | 0.05 | No data |
|  |  | $2(27-02-19)$ | 0.05 | 0.11 | 0.99 |
|  |  | $3(2-02-19)$ | 0.10 | 1.10 | 0.25 |
| 2 | S OO 18.603' E 100 09.901' | $1(27 / 2 / 2019)$ | 0.15 | 0.81 | 0.36 |
|  |  | $2(27-02-19)$ | 0.57 | 0.36 | 0.55 |
|  |  | $3(1-02-1019)$ | 0.64 | 0.56 | 0.23 |
| 3 | S 00 19.729' E 100 09.894' | $1(28-02-2019)$ | 0.56 | 0.94 | 1.78 |
|  |  | $2(1-3-2019)$ | 0.47 | 0.19 | 0.85 |
|  |  | $3(1-3-2019)$ | 0.56 | 0.38 | 1.69 |
| 4 | S 00 15.778' E 100 12.608' | $1(14-3-2019)$ | 0.40 | 0.68 | 0.18 |
|  |  | $2(15-3-2019)$ | 0.71 | 0.57 | 0.39 |
|  |  | $3(16-3-2019)$ | 1.63 | 1.12 | 1.47 |

Table E. Expected and observed presence and absence for field data validation of Rinuak.

| No | Long | Lat | Expected | Observed |
| :--- | :---: | :---: | :---: | :---: |
| 1 | 100.2204 | -0.3017 | 0 | 1 |
| 2 | 100.2204 | -0.3017 | 0 | 1 |
| 3 | 110.2239 | -0.2812 | 0 | 0 |
| 4 | 110.2239 | -0.2812 | 0 | 0 |
| 5 | 100.2132 | -0.2741 | 0 | 0 |
| 6 | 100.2132 | -0.2741 | 0 | 0 |
| 7 | 100.1992 | -0.2605 | 0 | 1 |
| 8 | 100.1992 | -0.2605 | 0 | 1 |
| 9 | 100.1931 | -0.2597 | 1 | 0 |
| 10 | 100.1931 | -0.2597 | 1 | 0 |
| 11 | 100.1867 | -0.2596 | 0 | 0 |
| 12 | 100.1867 | -0.2596 | 0 | 0 |
| 13 | 100.1783 | -0.2668 | 0 | 0 |
| 14 | 100.1783 | -0.2668 | 0 | 0 |
| 15 | 100.1706 | -0.2652 | 1 | 0 |
| 16 | 100.1706 | -0.2652 | 1 | 0 |
| 17 | 100.1667 | -0.273 | 1 | 0 |
| 18 | 100.1667 | -0.273 | 1 | 0 |
| 19 | 100.1637 | -0.2921 | 0 | 0 |
| 20 | 100.1637 | -0.2921 | 0 | 0 |
| 21 | 100.1683 | -0.3017 | 0 | 0 |
| 22 | 100.1683 | -0.3017 | 0 | 0 |
| 23 | 100.1721 | -0.3035 | 0 | 0 |
| 24 | 100.1721 | -0.3035 | 0 | 0 |
| 25 | 100.1763 | -0.3058 | 0 | 0 |
| 26 | 100.1763 | -0.3058 | 0 | 0 |
| 27 | 100.1686 | -0.3227 | 0 | 0 |
| 28 | 100.1686 | -0.3227 | 0 | 0 |
| 29 | 100.1854 | -0.3565 | 0 | 0 |
| 30 | 100.1854 | -0.3565 | 0 | 0 |
| 31 | 100.1951 | -0.3983 | 0 | 0 |
| 32 | 100.1951 | -0.3983 | 0.3979 | 0 |
| 33 | 100.1964 | -0.3979 | 0 | 0 |
| 34 | 100.1964 | 0.3918 | 0 | 0 |
| 35 | 100.1954 | 0.3918 | 0 | 0 |
| 36 | 100.1954 | 00.2174 | -0.3405 | 0 |
| 37 | 100.2174 | -0.3405 | 0 | 0 |
| 38 | 100.2213 | -0.3213 | 0 | 0 |
| 39 | 100.2213 | -0.3213 | 0 | 0 |
| 40 |  |  | 0 | 0 |
|  |  | 0 | 0 | 0 |
|  |  |  | 0 | 0 |

Table F. Expected and observed presence and absence for field data validation of Bada.

| No | Long | Lat | Expected | Observed |
| :--- | :---: | :---: | :---: | :---: |
| 1 | 100.2241 | -0.3028 | 1 | 1 |
| 2 | 100.2204 | -0.3017 | 1 | 1 |
| 3 | 100.2222 | -0.2997 | 1 | 1 |
| 4 | 100.2205 | -0.2973 | 1 | 1 |
| 5 | 110.2239 | -0.2812 | 1 | 1 |
| 6 | 100.2183 | -0.2801 | 0 | 1 |
| 7 | 100.2151 | -0.2789 | 0 | 0 |
| 8 | 100.2132 | -0.2741 | 0 | 0 |
| 9 | 100.2095 | -0.2719 | 0 | 0 |
| 10 | 100.1992 | -0.2605 | 1 | 1 |
| 11 | 100.1978 | -0.2611 | 1 | 1 |
| 12 | 100.1931 | -0.2597 | 1 | 1 |
| 13 | 100.1867 | -0.2596 | 1 | 1 |
| 14 | 100.183 | -0.2644 | 1 | 1 |
| 15 | 100.1783 | -0.2668 | 1 | 0 |
| 16 | 100.1706 | -0.2652 | 0 | 1 |
| 17 | 100.1667 | -0.273 | 1 | 0 |
| 18 | 100.176 | -0.2688 | 1 | 1 |
| 19 | 100.1637 | -0.2921 | 1 | 1 |
| 20 | 100.166 | -0.2991 | 1 | 1 |
| 21 | 100.1683 | -0.3017 | 1 | 1 |
| 22 | 100.1721 | -0.3035 | 1 | 1 |
| 23 | 100.1763 | -0.3058 | 1 | 1 |
| 24 | 100.1733 | -0.3119 | 1 | 1 |
| 25 | 100.1686 | -0.3227 | 1 | 1 |
| 26 | 100.1679 | -0.3309 | 1 | 1 |
| 27 | 100.1713 | -0.3336 | 1 | 1 |
| 28 | 100.1854 | -0.3565 | 0 | 0 |
| 29 | 100.1693 | -0.38 | 0 | 1 |
| 30 | 100.1695 | -0.3844 | 0 | 1 |
| 31 | 100.1729 | -0.3877 | 0 | 1 |
| 32 | 100.1815 | -0.3955 | 0 | 1 |
| 33 | 100.1881 | -0.3983 | 0 | 1 |
| 34 | 100.1951 | -0.3983 | 0 | 1 |
| 35 | 100.1964 | -0.3979 | 1 | 1 |
| 36 | 100.1954 | -0.3918 | 0.3405 | 0 |
| 37 | 100.2174 | -00.2196 | -0.3213 | 0.3019 |


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[^1]:    Note: Df: degree of freedom; AIC: Akaike information criteria; LL: log likelihood; LRT: likelihood ratio test against null model, LRTC: likelihood ratio test between models with habitat and sampling location as variables and models with habitat only.
    ${ }^{\text {a }}$ the best model

