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# Using species distribution models to optimize vector control in the framework of the tsetse eradication campaign in Senegal

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Tsetse flies are vectors of human and animal trypanosomoses in sub-Saharan Africa and are the target of the Pan African Tsetse and Trypanosomiasis Eradication Campaign (PATTEC). Glossina palpalis gambiensis (Diptera: Glossinidae) is a riverine species that is still present as an isolated metapopulation in the Niayes area of Senegal. It is targeted by a national eradication campaign combining a population reduction phase based on insecticide-treated targets (ITTs) and cattle and an eradication phase based on the sterile insect technique. In this study, we used species distribution models to optimize control operations. We compared the probability of the presence of G. p. gambiensis and habitat suitability using a regularized logistic regression and Maxent, respectively. Both models performed well, with an area under the curve of 0.89 and 0.92, respectively. Only the Maxent model predicted an expert-based classification of landscapes correctly. Maxent predictions were therefore used throughout the eradication campaign in the Niayes to make control operations more efficient in terms of deployment of ITTs, release density of sterile males, and location of monitoring traps used to assess program progress. We discuss how the models' results informed about the particular ecology of tsetse in the target area. Maxent predictions allowed optimizing efficiency and cost within our project, and might be useful for other tsetse control campaigns in the framework of the PATTEC and, more generally, other vector or insect pest control programs.

area-wide integrated pest management | genetic control | aerial release | chilled adult technique

setse are vectors of human African trypanosomosis, a major neglected tropical disease (1), and African animal trypanosomosis (AAT), one of the most important pathological constraints to livestock development in 38 infested African countries (2). The Pan African Tsetse and Trypanosomiasis Eradication Campaign is a political initiative started in 2001 that calls for intensified efforts to reduce the tsetse and trypanosomosis problem (3). As part of this global effort, the government of Senegal embarked in 2007 on a tsetse eradication campaign in a 1,000-km<sup>2</sup> target area of the Niayes region, neighboring the capital Dakar. In this area, the limits of the distribution of the tsetse target populations were assessed using a stratified entomological sampling frame based on remote sensing indicators (4). The only tsetse species present was Glossina palpalis gambiensis Vanderplank, and it was responsible for the cyclical transmission of three trypanosome species, namely Trypanosoma vivax, T. congolense, and T. brucei brucei, listed in order of importance (5). The high prevalence of animal trypanosomosis in the Niayes (serological prevalence of 28.7% for T. vivax) hampered peri-urban intensification of cattle production (particularly dairy cattle). A population genetics study demonstrated that the *G. p. gambiensis* population of the Niayes was completely isolated from the main tsetse belt in the southeastern part of Senegal (6). This comprehensive set of entomological, veterinary, genetic, and other baseline data confirmed the isolated nature of the *G. p. gambiensis* population in the Niayes, which prompted the government of Senegal to select an eradication strategy using area-wide integrated pest management (AW-IPM) principles (7).

The successful implementation of an AW-IPM strategy requires a thorough understanding of the ecology of the target population, particularly its spatial distribution, a study which was undertaken in the Niayes from 2007 to 2011 before the start of the operational eradication efforts. The selected strategy integrates insecticide-treated targets (ITTs) and cattle with the release of sterile insects. As the habitat of *G. p. gambiensis* is very fragmented (4), the targeting of suitable habitats for deployment of the ITTs is crucial to optimize cost efficiency (8) but also to enable the selection of appropriate sites for deployment of the

#### **Significance**

Tsetse flies transmit human and animal trypanosomoses in sub-Saharan Africa, respectively a major neglected disease and the most important constraint to cattle production in infested countries. They are the target of the Pan African Tsetse and Trypanosomiasis Eradication Campaign (PATTEC). Here we show how distribution models can be used to optimize a tsetse eradication campaign in Senegal. Our results allow a better understanding of the relationships between tsetse presence and various environmental parameters measured by remote sensing. Furthermore, we argue that the methodology developed should be integrated into future tsetse control efforts that are planned under the umbrella of the PATTEC initiative. The results have a generic value for vector and pest control campaigns, especially when eradication is contemplated.

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monitoring traps to assess the impact of the control campaign. Although the initial entomological sampling was well-developed and efficiently implemented in the target area (4), we deem the development and use of population distribution models to be very beneficial in this regard.

The use of species distribution models to optimize vector or pest control is quite novel. The existing tsetse distribution models (SI Text) were critical for a better understanding of tsetse distribution and AAT epidemiology, but their spatial resolution was not sufficient to guide an eradication process. In this paper, we used high-resolution images and recent advances in species distribution modeling methods to improve prediction accuracy. Predictive models, and more specifically machine-learning methods, were used to model the distribution of G. p. gambiensis (9). Model choice has an impact on the final output and also depends on the available data. In this study, both presence and absence data were available, which is uncommon with respect to tsetse data. Therefore, following Brotons et al., we used both datasets (10). Understanding how predictions from presence-absence models relate to predictions from presence-only models is important, because presence data are more reliable than absence data.

Presence–absence data were modeled using a regularized logistic regression to avoid overfitting with respect to model parameters. There are various approaches to regularization for least square methods in statistical learning. The most widely used are the ridge regression and the lasso. Ridge regression bounds the regression coefficient space by adding the L2 norm (root square of the sum of squared values) of the coefficients to the residual sum of squares, whereas the lasso is a penalized least square method that shrinks the coefficient space by imposing an L1 penalty (sum of absolute values) on the regression coefficients. The elastic net framework used here is a compromise that combines ridge regression (L2 penalization) and the lasso (L1 penalization) for more flexibility in model selection (11). This model was chosen because of its flexibility and capacity to penalize complex models (12).

A Maxent model was in addition used to model presence-only data (13). This model, which is one of the most widely used to model species distributions, is a machine-learning method based on maximum entropy. Absence data are replaced with so-called background data, which are a random sample of the available environment. Maxent fits a penalized maximum-likelihood model to avoid overfitting (L1 penalization). The logistic output from Maxent is a habitat suitability index rescaled to range from 0 to 1. Recently, the equivalence between Maxent and an infinitely weighted logistic regression was pointed out (14).

The relationship between occurrence and environmental data was explored with multidimensional exploratory analysis. For this purpose, we used the ecological factor niche analysis (ENFA) (15), which is a presence-only multidimensional method based on the concept of ecological niche (16).

The goal of this paper is to show how we selected among two competing approaches of species distribution modeling based on a large and accurate set of presence–absence data and how we used the results to optimize the eradication campaign in the Niayes of Senegal.

#### Results

**Ecological Niche.** All of the variables associated with forest habitats and mean normalized difference vegetation index (NDVI) were highly correlated with the presence of *G. p. gambiensis* (Fig. 1). Conversely, night and day land surface temperature (LST) ranges and maximum middle infrared (MIR) were negatively correlated with the presence of *G. p. gambiensis*. High values of these variables corresponded to lower vegetation cover, reducing the buffering of macroclimatic conditions by the vegetation.

Standard deviations of MIR, NDVI, and day and night LST had high values on the specialization axis (*SI Methods*), whereas the environmental envelope for *G. p. gambiensis* was very narrow on this axis. This suggested that these satellite-derived parameters



**Fig. 1.** First plan of the ecological niche factor analysis. The light gray polygon shows the overall environmental conditions available in the study area, the dark gray one shows environmental conditions where *G. p. gambiensis* was observed (representation of the realized niche), and the small white circle corresponds to the barycenter of its distribution. The first axis (marginality axis) is a measure capturing the dimension in the ecological space in which the average conditions where the species lives differ from the global conditions; a large marginality value implies that the conditions where the species is found are "far" from the global environmental conditions. In contrast, the second axis (specialization) measures the narrowness of the niche (ratio of the multidimensional variances of the available to occupied spaces). The eigenvalues of the axes present the percentages of the global inertia explained by theses axes (69% for the first plan). avg, average; mar, marginality; max, maximum; min, minimum; forest, surface of forest landscapes (Landsat data); spe, specialization; stddev, SD.

captured important environmental features for the species for which variations were poorly tolerated.

**Model Outputs.** Regularized logistic regression and Maxent presented similar receiver operating characteristic (ROC) curves using the presence–absence validation dataset, with areas under the curve (AUCs) of 0.89 and 0.92, respectively (Fig. 2). Their predictions were highly correlated (Fig. S1; r = 0.68, P < 0.01; Fig. 3). Maxent presented a slightly better sensitivity using 0.5 as a threshold for presence, or the threshold allowing the best percentage of correctly classified (PCC) index, and a better specificity when the threshold was set to enable a sensitivity of 0.96 (Table 1). Using the expert-based landscape classification derived from aerial photography, Maxent predicted suitable habitats better than the regularized logistic regression (AUC 0.79 and 0.58; Fig. 2).

The regularized training gain (likelihood-based measure of model quality) of the Maxent habitat suitability model is presented in Fig. S2. The Maxent models are presented using a single predictor or all of the predictors with the exception of the one of interest. The mean area of forest was the predictor giving the best gain when used alone, followed by the mean NDVI and the maximum forest area. These three predictors were highly correlated with each other along the ENFA marginality axis (Fig. 1 and *SI Methods*). Moreover, night LST SD had the highest negative effect on the gain (even if it was still very limited) when it was removed; thus, it contained specific information not accounted for by other predictors. The negative effect on the gain was the most prominent (6%) when all Landsat-related predictors (average, minimum, maximum, and SD of forest areas) were removed.



**Fig. 2.** ROC curves and AUCs of the regularized logistic regression and Maxent models. These results were obtained using the presence–absence validation dataset (*Left*) and the expert-based landscape classification from aerial images (*Right*).

Marginal response curves for the different predictors (Fig. S3) showed that the percentage of the area covered by forest was positively correlated with habitat suitability, with a sharp increase after 5% of forests and a plateau after 20%. With respect to average day LST, habitat suitability was stable between 20 and 35 °C and then sharply decreased; whereas average day LST was not low enough in the study area to limit the presence of G. p. gambiensis, high temperatures were unsuitable. The correlation between minimum day LST and habitat suitability was bell-shaped, with a maximum between 20 and 25 °C. This could be related to the buffering effect of dense tree cover on temperature drops at night. Indeed, night temperature drops faster in open environments than in the forest. Habitat suitability dropped sharply, with minimum temperatures exceeding 25 °C. The correlation between mean NDVI and habitat suitability was a sigmoid curve that increased sharply after 0.2. Regarding NDVI range, a plateau of habitat suitability was observed until 0.5 and sharply decreased after this threshold, thus illustrating the importance of perennial tree vegetation for this tsetse species. The relationship with habitat suitability was similar for MIR SD, related to the strong negative correlation between NDVI and MIR.

Use of Maxent Predictions in the Tsetse Eradication Project. Before the availability of the Maxent model in the Senegal project, operational choices such as selection of trap sites were made using a vegetation classification obtained from a Landsat 7 Enhanced Thematic Mapper Plus (ETM+) image of April 2001. Suitable habitat for *G. p. gambiensis* was mapped from this classification with high sensitivity but low specificity (4). The availability of the Maxent predictions was used in four ways to optimize the implementation of the eradication effort.

First, boundaries of the target eradication area, shown as black grid cells in Fig. 3, were validated by the model. All of the new suitable habitat areas identified by the Maxent model within a range of 5 km around sites where wild *G. p. gambiensis* had been trapped (at any occasion) were already included in the control strategy (*Discussion*). The target area was subdivided into four operational blocks, which are being sequentially addressed during the operational program (Fig. 4). Each block was subjected to a 1-y tsetse-density reduction phase, followed by an 18-mo eradication period using the sterile insect technique.

Second, the spatial distribution of the monitoring traps deployed since January 2012 in block 1 and since October 2012 in block 2 was modified through relocating 22 of the 97 monitoring traps (23%) to more suitable sites according to habitat suitability as predicted by the Maxent model (Fig. S4). In block 3, where monitoring had not started yet at the time of writing, the monitoring traps will also be deployed in sites that have a high (predicted) suitability value as indicated by the model.

Third, 1,347 insecticide-impregnated traps used for tsetse-density reduction were deployed in block 2 according to predictions of

Fourth, predicted suitable habitats were also used to optimize aerial releases of sterile male *G. p. gambiensis*. Polygons with similar surface areas of suitable patches were identified (RL1 and RL2 in block 1, for example; Fig. 4) and the density of released sterile males was adjusted proportionally to the area of suitable habitat in these polygons (0.24 and 0.11 flies per ha in RL1 and RL2, respectively).

**Optimization of Tsetse Suppression and Eradication by the Model.** In the first block, the apparent density of *G. p. gambiensis* dropped from an average of 0.42 (SD 0.39) flies per trap per d before control to an average of 0.04 (SD 0.11) flies per trap per d and 0.003 (SD 0.01) flies per trap per d during the suppression and eradication phases, respectively (Fig. 5). In the second block, the apparent density dropped from an average of 1.24 (SD 1.23) flies per trap per d during the suppression per trap per d before control to an average of 0.005 (SD 0.017) flies per trap per d during the suppression phase.

flies per trap per d during the suppression phase. Apparent fly density in block 2 was higher than in block 1 (generalized linear mixed model, P = 0.009; *SI Methods*) before the start of the suppression, which initially sharply reduced testse densities (P = 0.001). This effect was limited in time (6 mo), and tsetse density remained stable and even increased with time thereafter ( $P < 10^{-3}$ ), although remaining at a very low level (<0.04 fly per trap per d) until the eradication phase started. The cumulated reduction of densities with time was higher in block 2 (reduction of 99.6%) than in block 1 (reduction of 90.4%) ( $P < 10^{-3}$ ).

In block 1, the last wild fly was captured on August 9, 2012, ~6 mo after the start of sterile male releases. It was an old female (more than 40 d old) in its fourth larviposition cycle with an empty uterus, and the next follicle was immature and small, indicating an abortion. This female showed a copulation scar and its spermatheca was 85% filled, indicating that its sterility was probably induced by a mate with one sterile male. From the beginning of the eradication phase in block 1 (March 16, 2012) to the date corresponding to the last capture, only three other wild



**Fig. 3.** Predictions of the regularized logistic regression and Maxent models. In the case of the logistic model, the predictions correspond to probabilities of presence, whereas in the case of the Maxent model, they correspond to a rescaled suitability index (logistic output). The black cells correspond to the target area. Black and red crosses represent absence and presence data, respectively.

#### Table 1. Prediction qualities of the regularized logistic regression and Maxent models using the validation dataset

Model	Threshold	Sensitivity	Specificity	PCC	AUC
Glmnet	0.5	0.63 (0.07)	0.93 (0.03)	0.78 (0.04)	0.89 (0.03)
Maxent	0.5	0.64 (0.06)	0.96 (0.03)	0.80 (0.04)	0.92 (0.02)
Glmnet	0.11*	0.96 (0.03)	0.50 (0.07)	0.73 (0.04)	_
Maxent	0.13*	0.96 (0.03)	0.57 (0.07)	0.77 (0.04)	—
Glmnet	0.35 <sup>†</sup>	0.82 (0.05)	0.89 (0.04)	0.86 (0.03)	_
Maxent	0.32 <sup>†</sup>	0.86 (0.05)	0.79 (0.06)	0.82 (0.04)	_

These results were obtained with the presence-absence validation dataset. SDs are presented in parentheses. Glmnet, regularized logistic regression. \*Selected to optimize sensitivity.

<sup>†</sup>Selected to optimize PCC.

females could be dissected and all had indications of having mated with a sterile male. The average percentage of sterile males as a proportion of all catches was then 99.2% (SD 1.6%), corresponding to a sterile-to-wild male ratio of 130. The percentage of sterile males remained 100% thereafter (no wild fly was captured for 78 weekly collections with 25 monitoring traps), corresponding to a very likely eradication (probability of not detecting potential remaining flies of 0.002 only).

#### Discussion

Ecological Niche of G. p. gambiensis in the Niayes. Our analysis showed that the ecological niche of G. p. gambiensis in the Niayes area corresponded to permanent ligneous vegetation with a tree density sufficient to provide adequate shade and buffer temperature and relative hygrometry variations in comparison with macroclimatic conditions occurring in the surrounding open environments. Air temperature in dense tree vegetation in gallery forests can be 4 °C lower compared with the surroundings, and relative humidity 15% higher. This habitat provides resting sites for G. p. gambiensis in contrast to the more open habitat into which they may disperse for short periods (some hours) in search of a blood meal, namely their hunting sites. Suitable G. p. gambiensis habitat may be seasonal because of the variations of macroclimatic conditions and of nonpermanent vegetation.

The Maxent model confirmed that permanent dense tree vegetation was important for G. p. gambiensis, but also that the larger the area occupied by the flies during the rainy season (corresponding to their dispersal capacity) the more suitable this habitat appears to be for Ĝ. p. gambiensis. Evidence is provided by the positive correlation of the forest range with the ENFA marginality axis (Fig. 1).

The data layers derived from Landsat images, which have a higher spatial resolution (30 m) than Moderate Resolution

Imaging Spectroradiometer from the National Aeronautics and Space Administration (MODIS) data (250 m), were important predictors of the presence of G. p. gambiensis. This was expected, given their ability to survive in very small vegetation patches in the Niayes (4). It would be difficult to obtain such fine-scale data for larger regions of West Africa. Fortunately, a good correlation was found between forest predictors and minimum night LST (Fig. 1). Moreover, the regularized gain is only reduced by 6% when all forest predictors are removed.

#### Comparison of the Regularized Logistic Regression and Maxent Models.

The nature of predictions differs between regularized logistic regression (probability) and Maxent (index). However, the two predictions were highly correlated, as observed elsewhere (17). Model quality-assessment metrics were similar using the presence-absence validation dataset. Maxent predicted suitable areas better than regularized logistic regression based on the expert-based landscape classification. Tsetse presence data are generally more meaningful than absence data, because all known traps have a very low efficiency with respect to trapping rates (as a percentage of available individuals), namely  $\leq 1\%$  per d per km<sup>2</sup> (4). In the Niayes, observed trapping efficiencies were as low as 0.3% per d per km<sup>2</sup> following the release of more than 200,000 sterile male flies. In our case, particular conditions (feasibility study of an eradication project) allowed a rigorous selection of false absences (*SI Methods*), including homogeneous trapping protocol and known trap efficiency (4). Under other circumstances (trap efficiency unknown, different trap models, or trapping protocols), trapping may generate false absence data. Using only presence data to assess habitat suitability has the advantage that data derived from different sources (e.g., compilation of published data) can be combined to inform control projects.

Moreover, Maxent predictions were important for the eradication program as all suitable habitat needed to be included in the monitoring and the target area, even if they were not infested at the time of sampling due to possible movement among the patches. In the Niayes area, G. p. gambiensis are never present in all suitable patches at the same time. Instead, they form a metapopulation with patches connected through dispersal (18).

#### Use of Model Predictions for Optimization of the Eradication Project.

To generate a binary presence-absence map to delimitate the infested area, we chose a threshold providing a high sensitivity (96%). Indeed, in the case of an eradication project, it is paramount to reduce false negatives as much as possible, to avoid leaving tsetse-infested pockets not subjected to the control effort. Such areas can act as a source of reinvasion into previously cleared areas. During the entomological baseline data survey, the target area was divided into operational  $5 \times 5$  km grid cells where at least one tsetse had been captured, plus a buffer zone

> Fig. 4. Optimization of the integrated control strategy using model predictions. The Maxent model was used with a threshold of 0.13 to predict the suitable habitats for G. p. gambiensis (sensitivity of 0.96 and specificity of 0.57). In block 1, the suitable habitats allowed delimitating two polygons for aerial releases (RL1 and RL2) where the minimum numbers of sterile males released per km<sup>2</sup> were 24 and 11, respectively, based on Maxent predictions. Adult chilled tsetse flies are released with a Mubarqui smart release machine on board a gyrocopter of the Kalahari aerodrome (Upper Left) (20). In block 1, the green and gray lines represent the track flying records of the gyrocopter on April 14, 2014 in RL1 and April 11, 2014 in RL2, respectively. In block 2, 1,347 insecticide-impregnated traps were set from December 2012 to February 2013 in the predicted suitable sites (blue lozenges) to suppress tsetse populations.





**Fig. 5.** Impact of the control operations on tsetse apparent densities per trap per d. The red vertical lines represent the start of the suppression phase (insecticide targets) in each block, whereas the blue vertical lines represent the start of the eradication phase (release of sterile males). Densities are presented in natural log scale [log (catch + 1)]. The regression curves correspond to a nonparametric loess curve with associated 95% confidence interval. The map presents the location of the blocks that are targeted sequentially following a rolling carpet approach. The vertical red arrow shows the capture of the last indigenous fly in block 1.

consisting of the grid cells contiguous to these infested cells. This strategy was validated by the Maxent predictions using this sensitivity threshold and confirmed that the target area was spatially isolated from any other suitable area for *G. p. gambiensis*. The area selected for model predictions did not include the populations of northern Sine Saloum (4) because they represent a different ecotype, genetically isolated from the Niayes metapopulation. We however included the "small coast" area of Senegal, again to avoid leaving tsetse-infested pockets south of the target area (6).

Interestingly, the Maxent model predicted some suitable habitat north of the target area (Fig. 3), which was infested by *G. p. gambiensis* in the 1970s and was subjected to a control program using residual spraying of dieldrin and trapping (4). This former control program probably eliminated some pockets of *G. p. gambiensis* that were completely isolated from the main infested area by sand dunes. Actually, no *G. p. gambiensis* flies were captured in these sites despite intensive sampling for several months and using numerous traps (4), despite these sites appearing fully suitable for this species based on phytosociological criteria.

The improvement provided by the Maxent model, in comparison with the maps of suitable habitats (forests) based on the supervised classification of Landsat ETM+ images, is mainly related to specificity: 0.43 with this classification vs. 0.57 with Maxent. Sensitivity was already 0.96 with the supervised classification. Maxent selected those permanent tree habitats where climatic conditions (particularly temperature) allowed tsetse survival. Even if MODIS data mainly provide information on the macroclimate, they also allow making inferences on tree cover (particularly using night LST) and therefore on the buffering effect of permanent vegetation on macroclimatic conditions. Maxent models thus allowed increasing the sensitivity of the monitoring system, because traps previously set in unsuitable habitats have been moved based on suitability predictions (Fig. S4).

Regarding the suppression strategy, the use of the vegetation classification in block 1 (Maxent predictions were not available then) already represented a great improvement in comparison with previous tsetse control programs; 269 targets were set in block 1 from December 2010 to February 2011 using this classification (corresponding to 19.4 targets per km<sup>2</sup> of suitable habitat). They allowed a good suppression of the flies, whereas in the absence of a model of suitable habitat, densities of 60 targets per km<sup>2</sup> were required in Guinea against the same subspecies: Recently, on Fotoba Island, 30 targets per km<sup>2</sup> could only reduce the apparent fly density by 62% (8). In block 2, Maxent further improved the results, and a significantly higher reduction rate (99.6%) than in block 1 was obtained with a lower target density of 16.7 targets per km<sup>2</sup> of suitable habitat (and 2.7 targets per km<sup>2</sup> of the target area only). Considering a target cost of  $\in 3$  in our project and a deployment cost of  $\in 6$  per trap (19), the total reduction of the costs can be estimated at  $\sim \in 43,700$  for 1,000 km<sup>2</sup>.

Moreover, Maxent predictions allowed concentrating the release of sterile flies onto suitable habitats in block 1: With only 16.5 (SD 7.0) sterile flies released weekly per km<sup>2</sup>, a sterileto-wild male ratio of 130 was obtained, inducing 100% sterility in females and thus driving the population to extinction. The minimal weekly release rate was set at 10 and 100 flies per km<sup>2</sup> of unsuitable and suitable habitat, respectively. Initially, sterile males were released from the air using carton boxes dropped from a gyrocopter, but since February 2014 a more advanced automatic chilled adult release machine was used (Mubarqui smart release machine; Fig. 4) that can be parameterized daily considering these parameters and the amount of flies available at the emergence center (Senegalese Institute for Agricultural Research) (20, 21). Within this machine, the vibratory feeders can be adjusted to release rates between 10 and 100 flies per km<sup>2</sup>, with a gyrocopter flying at a speed of 110 km/h. In the absence of the Maxent models, it would have been necessary to release at least 100 flies per km<sup>2</sup> everywhere in the target area [in Zanzibar, up to 300 flies per km<sup>2</sup> were even released in the forest section to eradicate G. austeni (22)]. With a mean cost of  $\in 0.2$ per pupa [including production costs at Centre International de Recherche-Développement sur l'Elevage en zone Subhumide (CIRDES), Burkina Faso and the Slovak Academy of Sciences (SAS) and  $\in 0.04$  transport cost per pupa], this represents major savings of ~€590,000 for our project, taking into account that 70% of received pupae provide for operational sterile flies, when taking into account the emergence rate and mortality rate at the insectarium before release. Moreover, the total number of sterile male G. p. gambiensis pupae produced by the CIRDES and the SAS is currently limited to 25,000 pupae per wk, and treating the full area would not have been possible without adjusting release densities to the availability of suitable habitat.

The same methodology will be applied in the two remaining blocks following the rolling carpet approach: When eradication is started in a given block, suppression is started in the contiguous block to avoid any risk of reinvasion. Based on this strategy, the full target area is planned to be cleared from tsetse by the end of 2016.

The same approach might be used to optimize any vector or insect pest control program, especially when eradication is the selected strategy (23).

#### Methods

**Site Description.** The study area had a total surface of 7,150 km<sup>2</sup>, located in the Niayes region of Senegal (14.1° to 15.3° N and 16.6° to 17.5° W). At the time of the study, it was the target of an eradication campaign against an isolated *G. p. gambiensis* population (4) (Fig. 4). The climate is hot and dry from April to June, whereas the rainy season occurs from July to October and the cold dry season from November to March.

**Response Data.** *Learning dataset.* A cross-sectional survey was implemented from December 2007 to March 2008 (dry season) to collect baseline data for the eradication campaign using 683 unbaited Vavoua traps (4). Traps were

APPLIED BIOLOGICAL SCIENCES removed as soon as a tsetse fly was caught (minimum 1 d). In the absence of capture, the traps were retrieved after a maximum of 42 d.

To homogenize data quality and avoid pseudoreplication (data points in the same pixel), the dataset was simplified in two ways. First, the study area was rasterized into square pixels with a 250-m resolution to match the model predictors (*SI Methods*).

For the presence data, all trap positions located within the same pixel were aggregated and concentrated in the pixel center. Presence data were observed in 68 pixels from 91 presence points (Fig. S5).

For the absence data, all trap positions located within a buffer of 500 m around a presence point were removed (to account for tsetse dispersal capacity). Furthermore, we removed absence pixels with P > 0.01 that the flies were present despite the absence of trapping using the model described (24) (*SI Methods*). From the initial 592 absence data, 333 were finally retained (Fig. S5), 56 of which were used in the validation dataset. Therefore, the training dataset was composed of 68 presence and 269 absence data.

Validation datasets. Trap data were collected independent of the training dataset from April 2009 to February 2013 during different surveys (tsetse dispersal and competitiveness studies, longitudinal monitoring of the demographic structure of tsetse populations, etc.). This dataset included 92 presence and 64 absence data. It was processed as described above, resulting in a dataset of 64 presence and 1 absence data (trapping times were not long enough to ascertain tsetse absence). A subset of 56 absence data was extracted from the 333 absence data described above.

A second validation dataset was created using 182 aerial photos taken from a gyrocopter at an altitude ranging from 100 to 300 m. The environment identified from these pictures was subsequently categorized as suitable or unsuitable for tsetse habitat (4). Picture coordinates were corrected using Google Earth to take the angle and deviation from the ground into account. Overall, 23 suitable and 159 unsuitable habitats were identified, thereafter called "expert-based habitats" (Fig. S6).

**Predictors Used for Tsetse Suitable Habitat.** Climatic and environmental data were derived from a time series of MODIS (version V005). Composite day and night land surface temperature (MOD11A2; 8-d averages), middle infrared (MOD13Q1; 16-d averages), and normalized difference vegetation index (MOD13Q1; 16-d averages) were selected for the analysis (25). Spatial resolution of pixels was 250 × 250 m (*SI Methods*).

A supervised classification of the vegetation was achieved using Landsat 5 Thematic Mapper satellite images with a spatial resolution of  $30 \times 30$  m (*SI Methods*). Four cloud-free and haze-free satellite images were used, from

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October 2009, April 2010, June 2010, and December 2010, to take into account the seasonal dynamics of these habitats, named "forests" in the manuscript.

### **Models.** Ecological niche factor analysis was used to characterize the habitat of *G. p. gambiensis* in the target area (*SI Methods*).

Regularized logistic model was used to predict tsetse presence probability. Maxent analysis was used to predict habitat suitability. Approximately 10,000 pixels were sampled from the environmental variables (background) to calibrate the model. Because of the small sample size for presence data (56 points), only linear and quadratic transformations of environmental variables were used.

Model performance and comparison were assessed using the two validation sets (26). Model performance metrics were computed for each predictive model (27). The metrics used were the area under the receiver operating curve, called the area under the curve (28), the percentage of consonants correct (PCC), the specificity, and the sensibility (*SI Methods*).

**Regularized gain is an additional metric for Maxent models.** Marginal response curves were also computed to assess the relationship between the predicted suitability index and a given environmental datum. These curves were obtained by varying this variable while keeping all others at their average value.

Predicted values from regularized logistic regression and Maxent models were compared using Spearman's rank correlation coefficient.

**Impact of Control Operations.** We used a generalized linear mixed model (29) to measure the impact of the suppression on tsetse apparent densities. The response data were tsetse counts in the traps. Time (measured in weeks), treatment (suppression or not), and the block (1 or 2) and their interactions were used as fixed effects, whereas the trap locations were used as random effects (*SI Methods*). Raw data are presented in Dataset S1.

The probability that eradication was effective in block 1 was estimated using the same model used to clean the absence dataset (24) (*SI Methods*), considering that at least a couple of flies were necessary to maintain the population.

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