

# ForestAtRisk: A Python package for spatial modelling and forecasting of tropical deforestation

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# Summary

The ForestAtRisk Python package can be used to model the tropical deforestation spatially and forecast the future forest cover in the tropics. It includes functions to (i) sample the forest cover change observations and retrieve the information on potential spatial explanatory variables for each observation, (ii) estimate the parameters of various spatial deforestation models, (iii) predict the spatial probability of deforestation, (iv) forecast the likely forest cover in the future, (v) validate the models and the projected maps of forest cover change, and (vi) plot the results. Spatial information is provided through georeferenced raster files which can be very large (several gigabytes). Functions in the ForestAtRisk package treat large rasters by blocks of data, making computations fast and efficient. This allows analysis on large geographical extent (e.g. country scale), and at high resolution (e.g. 30 m). The ForestAtRisk package offers the possibility to use a logistic regression model with auto-correlated spatial random effects to model the deforestation process. Spatial random effects allow to structure the residual spatial variability (which is often large) within the model. They account for unmeasured or unmeasurable variables that explain a part of the residual spatial variation in the deforestation process that is not explained by the model's explanatory variables. In addition to bringing new features, the ForestAtRisk Python package is open source (license GPLv3), cross-platform and scriptable (through Python), user-friendly (functions provided with full documentation and examples), and easily extendable (with additional statistical models for example). The ForestAtRisk Python package has been recently used to model the deforestation and forecast the future forest cover by 2100 in the whole humid tropics.

### Statement of Need

The ForestAtRisk Python package can be used to model the tropical deforestation spatially and forecast the future forest cover in the tropics (Figure 1).

Functions in the ForestAtRisk package treat large rasters by blocks of data, making computation fast and efficient (with low memory usage).

Using functions from the ForestAtRisk Python package makes computation fast and efficient (with low memory usage) by treating large raster data by blocks. Numerical computations on blocks of data are performed with the NumPy (https://numpy.org) Python module whose core is mostly made of optimized and compiled C code which runs fast (Harris et al., 2020).

Using simple "glm" model is a commonly proposed approach for spatial modelling of deforestation (Eastman & Toledano, 2017; Ludeke et al., 1990; Mas et al., 2007; Mertens & Lambin, 1997; Rosa et al., 2014; B. S. Soares-Filho et al., 2002; Britaldo S. Soares-Filho et al., 2001; Verburg et al., 2002-09-21). The random forest model (Breiman, 2001) is a machine learning approach using an ensemble of random classification trees (where both observations and features are chosen at random to build the classification trees) to predict the deforestation

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#### Software

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probability for a forest pixel. Random forest has been intensively used for species distribution modelling (Thuiller et al., 2009) and is now also commonly used for spatial modelling of deforestation (Grinand et al., 2020; Santos, 2019; Zanella et al., 2017). The "glm" and "rf" models were fitted using functions LinearRegression and RandomForestClassifier respectively, both available in the scikit-learn Python package (Pedregosa et al., 2011).

Using observations of forest cover change in the period 2010-2020, we modelled the spatial probability of deforestation as a function of the n explanatory variables using a logistic regression. We considered the random variable  $y_i$  which takes value 1 if the forest pixel i was deforested in the period 2010–2020 and 0 if it was not. We assumed that  $y_i$  follows a Bernoulli distribution of parameter  $\theta_i$  (Eq. @ref(eq:icar)). In our model,  $\theta_i$  represents the spatial relative probability of deforestation for pixel i. We assumed that  $\theta_i$  is linked, through a logit function, to a linear combination of the explanatory variables  $X_i\beta$ , where  $X_i$  is the vector of explanatory variables for pixel i, and  $\beta$  is the vector of effects  $[\beta_1, \dots, \beta_n]$  associated to the n variables. All the continuous explanatory variables were normalized before fitting the model. The model includes an intercept  $\alpha$ . To account for the residual spatial variation in the deforestation process, we included an additional random effect  $ho_{j(i)}$  for each spatial cell j of a 10 imes 10 km grid covering each study-area. This grid resolution was chosen in order to have a reasonable balance between a good representation of the spatial variability of the deforestation process and a limited number of parameters to estimate. A sampled forest pixel i was associated to one cell j and one random effect  $\rho_{j(i)}$ . We assumed that random effects were spatially autocorrelated through an intrinsic conditional autoregressive (iCAR) model (Banerjee et al., 2014; Besag et al., 1991). This model is denoted "icar" in subsequent sections and results. In an iCAR model, the random effect  $\rho_j$  associated to cell j depends on the values of the random effects  $\rho_{j'}$  associated to neighbouring cells j'. In our case, the neighbouring cells are connected to the target cell j through a common border or corner (cells defined by the "king move" in chess). The variance of the spatial random effects  $\rho_i$  was denoted  $V_o$ . The number of neighbouring cells for cell j, which might vary, was denoted  $n_i$ . Spatial random effects  $\rho_i$ account for unmeasured or unmeasurable variables (Clark, 2005) that explain a part of the residual spatial variation in the deforestation process that is not explained by the fixed spatial explanatory variables  $(X_i)$ .

$$y_{i} \sim \mathcal{B}ernoulli(\theta_{i})$$

$$\log \mathsf{it}(\theta_{i}) = \alpha + X_{i}\beta + \rho_{j(i)}$$

$$\rho_{j(i)} \sim \mathcal{N}ormal(\sum_{j'} \rho_{j'}/n_{j}, V_{\rho}/n_{j})$$

$$(1)$$

Parameter inference is done in a hierarchical Bayesian framework. Function model\_bino mial\_iCAR() from the forestatrisk Python package was used for parameter inference. This function calls an adaptive Metropolis-within-Gibbs algorithm (Rosenthal & others, 2011) written in C for maximum computation speed.



# **Figures**

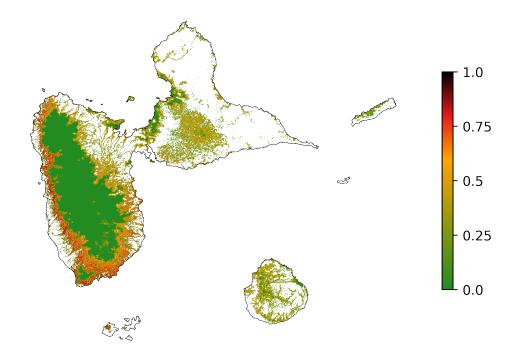


Figure 1: Map of the spatial probability of deforestation in the Guadeloupe archipelago. Colored pixels represent the extent of the natural old-growth tropical moist forest in 2020. A relative spatial probability of deforestation is computed for each forest pixel. The original map has a 30 m resolution. Probability of deforestation is a function of several explanatory variables describing: topography (altitude and slope), accessibility (distances to nearest road, town, and river), forest landscape (distance to forest edge), deforestation history (distance to past deforestation), and land conservation status (presence of a protected area). This map can be reproduced following the Get Started tutorial at <a href="https://ecology.ghislainv.fr/forestatrisk">https://ecology.ghislainv.fr/forestatrisk</a>.

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