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# Economics of land use reveals a selection bias in tree species distribution models

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# 28**Abstract**

- Aims. In human dominated ecosystems, the presence of a given species is the result of both 30the ecological suitability of the site and human impacts such as land-use choices. The influence of 31land-use choices on the predictions of species distribution models (SDMs) has however been often 32neglected. Here, we provide a theoretical analysis of the land-use selection bias affecting classical 33SDMs in the case of either presence-only or presence-absence datasets. Land-use selection bias in 34SDM predictions is then quantified for four widespread European tree species, using NFI data.
- 35 **Location**. Continental France
- Methods. We describe a Bivariate Selection Model (BSM) that estimates simultaneously the 37economics of land-use choices and species responses to bioclimatic variables. The land-use equa-38tion, based on an econometric model of landowner choices, is joined to an equation of species re-39sponse to bioclimatic variables.
- Results. We evidenced a significant land-use selection bias in all the species studied. The 41sign and the magnitude of the bias varied among species and were strongly related to the type of 42dataset used in the SDM calibration (presence-only or presence-absence). In addition, the BSM esti-43mate the spatial covariance between the probability of presence and the presence of compatible 44land-use. We found that, depending on the species, sites with high ecological suitability could 45present a high probability of compatible land-use (positive covariance) or a low probability (nega-46tive covariance).
- Main Conclusion. We showed that using classical SDMs in human dominated areas can 48lead to strong miss-estimations of actual species' distributions and could therefore prevents sound 49projections of the effect of climate change. Beyond, the proposed BSM represents a crucial step to 50account for economic forces shaping species distribution in anthropized area and pave the way for a 51direct assessment of trade-offs and opportunities that may arise in a context of global change.

# 521. Introduction

Species distribution models (SDMs) have been increasingly used in a wide range of 53 54ecological studies, with the general aim of understanding and predicting the distributional area of 55species (Guisan & Thuiller 2005; Thuiller et al. 2005; Zimmermann et al. 2010; Engler et al. 2011; 56Franklin *et al.* 2013). The data used to calibrate or evaluate SDMs often consist in presence-only (P-57O) data, i.e., a collection of point locations where a species is found, as absence data are difficult to 58obtain and require prohibitively expensive sampling designs. This constraint on data resources has 59been circumvented by using information on the range of the available environment conditions in the 60studied region (referred to as background data) instead of recorded absences (Elith et al. 2006). 61However, a fundamental limitation of SDM calibration based on P-O data is its sensitivity to the 62sample selection bias (Phillips et al. 2009) that can occur if the observations are not randomly 63distributed over a landscape. For instance predictions are likely to be affected by a sample selection 64bias if occurrences are preferably recorded near roads or towns (observer bias, Warton et al. 2013), 65or if sites expected to be suitable for a given species are over sampled (Jiménez-Valverde 2012; 66Smith 2013). In these cases, there is a risk that the recorded occurrences only partially describe the 67suitable species area of the studied landscape, which lead to wrong SDM predictions.

Because the sample selection bias affects both presence and absence records, its effect is 69usually thought to cancel out in the predictions of SDMs calibrated using presence-absence (P-A) 70data (Elith *et al.* 2011). But SDM predictions based on P-A data remain actually confronted to the 71general issue of selection bias, due to the uncertain nature of absence data (Jiménez-Valverde *et al.* 722008; Lobo *et al.* 2010; Duputié *et al.* 2014). Indeed, the absence of a species from any location 73cannot unambiguously be related to unsuitable environmental conditions as it can also result from 74internal processes related to the species ecology (biotic interactions, limited dispersion, habitat 75fragmentation) or from external processes driven by human activities (e.g., land-use choices, 76hunting intensity, resource management). In these cases, selection bias cannot be reduced by

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77improving the sampling scheme or increasing the number of sampled sites of presences and/or 78absences. The biases related to the imperfect detection of potential distribution are therefore very 79common in all the species distribution studies, even in the case of sessile species such as plants 80(Chen *et al.* 2013) and could strongly impact SDM performances (Lahoz-Monfort *et al.* 2014).

- The effect of land-use in SDM predictions has long been acknowledged. Indeed, SDMs are 82usually evaluated at continental scale and coarse resolution, at which climatic tolerance is a primary 83factor (Pearson & Dawson 2003; Thuiller *et al.* 2004). However, the degradation of habitats due to 84land-use changes remains a major cause of biodiversity loss (Bellard *et al.* 2012) and land-use 85choices could exert a strong control over species distribution evaluated at medium to fine spatial 86scales (1–10 km resolution, Pearson & Dawson 2003). Models integrating climate and land-use 87change components have been developed to produce sound projections of the future of species 88distribution (Brook *et al.* 2008). A few studies have integrated static (Pearson *et al.* 2004; Thuiller 89*et al.* 2004), or dynamic (Stanton *et al.* 2012; Martin *et al.* 2013) land-use variables along with 90climate variables in SDMs. All these studies came to the general conclusion that land-use effects on 91species distribution are largely over-ride by climatic effects. However, despite its potential 92importance, the effect of the selection bias created by land-use choices on SDM predictions has 93never been evaluated.
- In this study, we use an econometric model of land-use choices to correct the land-use 95selection bias that may affect tree species response to bioclimatic variables in classical SDMs. We 96provide a joint ecological and economical framework for modelling tree species distribution, taking 97into account the potential observed and unobserved selection bias induced by anthropogenic 98land-use choices. We first provide a theoretical analysis of the land-use selection bias in SDMs, 99generalizing the two classical SDM practices of P-O and P-A calibrations. Our conceptual approach 100also brings new insights into the ecological meaning of results obtained from both P-O and P-A tree 101distribution models in terms of effective and potential distributions (Jiménez-Valverde *et al.* 2008;

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102Lobo *et al.* 2010). Secondly, we estimate Bivariate Selection Models (BSMs) using empirical data 103from the French national forest inventory. In addition to the classically used environmental 104predictors (climate, soil and topography), BSMs draws on economic predictors (returns from crops, 105forests and urban uses). Thirdly, we compared the BSM predictions with those classical SDMs 106calibrated with P-O and P-A datasets, for four widespread European tree species: sessile oak 107(*Quercus petraea*), pubescent oak (*Quercus pubscens*), common beech (*Fagus sylvatica*) and silver 108fir (*Abies alba*) across a range of spatial resolutions (2 km, 4 km and 8 km). BSM simulations are 109shown to increase the interpretative content of niche modelling by (i) distinguishing between 110potential and effective probability of presence and (ii) explicitly estimating the spatial interactions 111between the ecological suitability for a tree species and the economical conditions of having a 112forested land-use.

# **2. Methods**

## **2.1.** General framework

We model the potential presence of a particular species through the continuous latent (i.e., 116not directly observed) variable  $\mu$ . Environmental conditions of a site i are potentially suitable 117for this species if the corresponding value  $\mu_i$  is greater than a given threshold, normalized to 0. 118On such sites with  $\mu_i > 0$ , the species of interest can be potentially observed. This latent variable 119is parametrized by an unknown smooth function  $f_p$ , a set of abiotic environmental variables 120 X and an additive random component  $\varepsilon$ , with  $\mu_i = f_p(X_i) - \varepsilon_i$ . The subscript p marks 121terms relative to the potential distribution. By noting  $(m_p = 1 \mid X_i)$  the event of "potentially 1220bserving the species of interest in the environment  $X_i$ ,", we have:

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$$\operatorname{Prob}\left(m_{p}=1 \mid X_{i}\right) = \operatorname{Prob}\left(\mu_{i}>0\right) = \operatorname{Prob}\left(\varepsilon_{i} < f_{p}\left(X_{i}\right)\right)$$
 (1)

However, in human-dominated ecosystems, the observation of a forest tree species is also 127strictly conditioned by the presence of a compatible land-use, i.e., the presence of forest rather than 128agricultural or urban areas. As it is common in the econometric literature concerning discrete 129individual choices, we modelled land-use choices using the random utility framework (Lubowski *et* 130*al.* 2008; Lewis 2010; Ay *et al.* 2014). Accordingly, we assumed that observed land-use is the result 131of utility maximization. Utility is specified as the sum of monetary returns a landowner can expect 132from land-uses and a random term representing the unobserved part of utility. This unobserved part 133potentially accounts for non-measurable variables or non-monetary values.

We assumed that the forested land-use is the only one compatible with the observation of 135 forest tree species. Considering that tree species inventories are typically restricted to forests, this 136 assumption is implicitly made by many other empirical studies using such data. We can then note 137  $\tilde{u}_i$  the difference between the utility from forest and the maximum utility from any other land-138 uses, which are incompatible with the presence of forest tree species. Hence, the compatible forest 139 land-use is chosen by landowners when this utility difference is strictly positive. This utility 140 difference is parametrized with a unknown smooth function  $f_i$ , the abiotic environmental 141 variables X, the monetary returns W expected by landowner, and an additive random 142 component  $\xi$ ,  $\tilde{u}_i = f_i(X_i, W_i) - \xi_i$ . The subscript I marks terms relative to land-use choices. 143 By noting  $(m_i = 1 \mid X_i, W_i)$  the event of "having a compatible land-use in the environment  $X_i$  144 with the economic conditions  $W_i$ ,", we note:

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$$\operatorname{Prob}(m_l = 1 | X_i, W_i) = \operatorname{Prob}(\tilde{u}_i > 0) = \operatorname{Prob}(\xi_i < f_l(X_i, W_i))$$
 (2)

#### 1482.2. Land-use selection bias

In the following, we develop a formal description of the land-use selection bias by 150modelling jointly tree species distribution and land-use choices. Note that the two previous events 151(i.e., equations (1) and (2)) require observations of very different nature to be estimated. 152Realizations of  $(m_p=1|X_i)$  are theoretical (i.e., related to potential presence) and are not 153directly observable from non-experimental data. Conversely, realizations of the land-use event 154  $(m_i=1|X_i,W_i)$  can be directly observed from any land-cover dataset. The land-use selection 155bias occurs because classical tree distributional data only contains the intersection of these two 156events, namely effective presence. As a consequence, what is classically observed and recorded in 157species distribution datasets is the event of "effectively observe the species in the environment 158  $X_i$  with the economic conditions  $W_i$ ", that we note  $(m_e=1|X_i,W_i)$ . The subscript e 159marks terms relative to the effective distribution that takes into account the land-use constraint.

Thus, the observable value  $m_e$  depends on the observable value  $m_l$  and the 161unobservable value  $m_p$ :

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163 
$$m_e = m_p \times m_l(3)$$

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165Eqn 3 illustrates the uncertain nature of effective species absences (i.e., the event  $m_e$ =0 ) which 166is potentially attributable to  $m_l$ =0 and/or  $m_p$ =0. Consequently,  $m_p$  is not directly 167deductible from  $m_e$ . Additional information on land-use compatibility has the potential to clarify 168the link between  $m_e$  and  $m_p$  events in classical SDM modelling.

The land-use selection bias arising from confounding  $m_e$  and  $m_p$  can be expressed 170differently for SDM calibrated with presence-only (P-O) and presence-absence (P-A) data, both 171being currently used in tree species distribution studies (e.g. Dorazio 2014; Fithian *et al.*, 2014 for

172P-O; e.g. Cheaib *et al* 2012; Gritti *et al*, 2013 for P-A). We choose to define P-A dataset as the case 173where absence records are limited to a compatible land use (i.e., forested sites where  $m_l$ =1 ). The 174absences recorded in the P-A dataset therefore only correspond to sites with unsuitable ecological 175conditions. This allows providing two contrasted benchmarked case study for modelers with clear 176mathematical results.

Predictions obtained from our P-O dataset correspond to the case where all sites are used in 178the SDM calibration (i.e., including sites with all types of land-use). This is equivalent to the typical 179case of occurrence records combined with naïve background data (e.g., Elith *et al.* 2006). The 180predictions obtained from our P-A dataset correspond to the case where only sites with a compatible 181land-use are used in the calibration (i.e., forested sites where  $m_l$ =1 ). The absences recorded in 182the P-A dataset therefore only correspond to sites with unsuitable ecological conditions, i.e. 183"confirmed" absences.

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Depending on whether P-O or P-A data are used for the calibration, classical SDMs 186respectively estimate one of the following probabilities instead of the probability of interest Prob  $(m_p=1 \mid X_i)$ :

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189 (P-O) 
$$\text{Prob}(m_e=1 | X_i, W_i) \text{ or } (\text{P-A}) \text{ Prob}(m_e=1 | X_i, m_i=1)$$
 (4)

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191The (P-O) case is the probability of the intersection of the events  $m_p=1$  and  $m_l=1$  for all 192sites:

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194 Prob 
$$(m_e=1 \mid X_i, W_i) = \text{Prob}(\varepsilon_i < f_p(X_i) \cap \xi_i < f_l(X_i, W_i))$$
 (5)

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196Note that, as the events  $m_l=1$  and  $m_l=0$  are a partition of the probability space (i.e., their

197probabilities sum to one), the probability of interest is the sum of this (P-O) probability and the 198probability of potential presence for incompatible land-uses:

200 
$$\operatorname{Prob}(m_p = 1 \mid X_i, W_i) = \operatorname{Prob}(\varepsilon_i < f_p(X_i) \cap \xi_i < f_l(X_i, W_i))$$

201 +Prob
$$\left(\varepsilon_{i} < f_{p}(X_{i}) \cap \xi_{i} \square f_{l}(X_{i}, W_{i})\right)$$
 (6)

203Consequently, we can express the bias from (P-O), i.e., the difference between the (P-O) probability 204and the probability of interest, as:

206 (P-O) bias = 
$$\operatorname{Prob}(m_e = 1 \mid X_i, W_i) - \operatorname{Prob}(m_p = 1 \mid X_i)(7) = -\operatorname{Prob}(\varepsilon_i < f_p(X_i) \cap \xi_i \square f_l(X_i, W_i))(8)$$

208Next, according to Bayes' theorem, the probability of interest can be expressed as:

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$$\operatorname{Prob}(m_p=1|X_i,W_i) = \operatorname{Prob}(m_p=1|X_i,m_l=1) \times \operatorname{Prob}(m_l=1|X_i,W_i)$$
. (9)

212Note that, if  $m_l=1$  , the events  $m_p=1$  and  $m_e=1$  are equivalent. So the P-A probability of 213equation (4) is similar to the term  $\operatorname{Prob}(m_p=1|X_i,m_l=1)$  from equation (9), and the associated 214P-A bias can be written as:

$$\mathbf{216} \quad \mathbf{P-A \ bias} = \operatorname{Prob}\left(m_{e} = \mathbf{1} \mid X_{i}, m_{l} = 1\right) - \operatorname{Prob}\left(m_{p} = \mathbf{1} \mid X_{i}\right) (\mathbf{10}) = \frac{\operatorname{Prob}\left(\varepsilon_{i} < f_{p}\left(X_{i}\right) \cap \xi_{i} < f_{l}\left(X_{i}, W_{i}\right)\right)}{\operatorname{Prob}\left(\xi_{i} < f_{l}\left(X_{i}, W_{i}\right)\right)} - \operatorname{Prob}\left(\varepsilon_{i} < f_{p}\left(X_{i}\right)\right) (\mathbf{10}) = \frac{\operatorname{Prob}\left(\varepsilon_{i} < f_{p}\left(X_{i}\right) \cap \xi_{i} < f_{l}\left(X_{i}, W_{i}\right)\right)}{\operatorname{Prob}\left(\xi_{i} < f_{l}\left(X_{i}, W_{i}\right)\right)} - \operatorname{Prob}\left(\varepsilon_{i} < f_{p}\left(X_{i}\right)\right) (\mathbf{10}) = \frac{\operatorname{Prob}\left(\varepsilon_{i} < f_{p}\left(X_{i}\right) \cap \xi_{i} < f_{l}\left(X_{i}, W_{i}\right)\right)}{\operatorname{Prob}\left(\varepsilon_{i} < f_{p}\left(X_{i}\right) \cap \xi_{i} < f_{p}\left(X_{i}\right)\right)} - \operatorname{Prob}\left(\varepsilon_{i} < f_{p}\left(X_{i}\right) \cap \xi_{i} < f_{p}\left(X_{i}\right)\right) = \frac{\operatorname{Prob}\left(\varepsilon_{i} < f_{p}\left(X_{i}\right) \cap \xi_{i} < f_{p}\left(X_{i}\right)\right)}{\operatorname{Prob}\left(\varepsilon_{i} < f_{p}\left(X_{i}\right) \cap \xi_{i} < f_{p}\left(X_{i}\right)\right)} - \operatorname{Prob}\left(\varepsilon_{i} < f_{p}\left(X_{i}\right) \cap \xi_{i} < f_{p}\left(X_{i}\right)\right) = \frac{\operatorname{Prob}\left(\varepsilon_{i} < f_{p}\left(X_{i}\right) \cap \xi_{i} < f_{p}\left(X_{i}\right)\right)}{\operatorname{Prob}\left(\varepsilon_{i} < f_{p}\left(X_{i}\right) \cap \xi_{i} < f_{p}\left(X_{i}\right)\right)} - \operatorname{Prob}\left(\varepsilon_{i} < f_{p}\left(X_{i}\right) \cap \xi_{i} < f_{p}\left(X_{i}\right)\right) = \frac{\operatorname{Prob}\left(\varepsilon_{i} < f_{p}\left(X_{i}\right) \cap \xi_{i} < f_{p}\left(X_{i}\right)\right)}{\operatorname{Prob}\left(\varepsilon_{i} < f_{p}\left(X_{i}\right) \cap \xi_{i} < f_{p}\left(X_{i}\right)\right)} - \operatorname{Prob}\left(\varepsilon_{i} < f_{p}\left(X_{i}\right) \cap \xi_{i} < f_{p}\left(X_{i}\right)\right) = \frac{\operatorname{Prob}\left(\varepsilon_{i} < f_{p}\left(X_{i}\right) \cap \xi_{i} < f_{p}\left(X_{i}\right)\right)}{\operatorname{Prob}\left(\varepsilon_{i} < f_{p}\left(X_{i}\right) \cap \xi_{i} < f_{p}\left(X_{i}\right)\right)} - \operatorname{Prob}\left(\varepsilon_{i} < f_{p}\left(X_{i}\right) \cap \xi_{i} < f_{p}\left(X_{i}\right)\right)$$

The eqn (8) and (11) allow us to determine unambiguously the directions of the bias that 219affects classical SDM modelling. In case of P-O calibration (eqn 8) the bias is negative, so classical 220P-O SDMs always under-estimate the probability of potential presence. Moreover, it appears from 221eqn (8) that the P-O bias depends on the intersection of two inequalities on errors, meaning that it 222depends on the correlations between the errors of the ecological and economical equations. More 223precisely, the bias decreases as the coefficient value of the correlation between the errors increases 224from -1 to 1 (see the formal proof in section 1.2 of Supporting Information, SI). 225To interpret the bias that affects P-A SDMs (equation 10-11) we should recall that, for 226independently distributed variables, the probability of the intersection is the product of the marginal 227probabilities. So, one can show by simplifying the numerator that the P-A bias is zero if the errors 228are independent. For positively correlated variables the probability of the intersection is greater that 229the product of marginal probabilities. In this case, the bias is positive and classical P-A SDMs over-

230estimate the probability of potential presence (see Section 1.2 of SI). In the case of negatively

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## 2332.3. Empirical Implementation

231correlated errors, this P-A bias is negative.

- We applied this theoretical framework on tree species distribution modelling by estimating 235Bivariate Selection Models (BSMs, Marra et al. 2014) that included both ecological and economical 236drivers. BSMs were estimated with the package "SemiParBIVProbit" available on CRAN (Marra et 237al. 2014). BSMs require exclusion restrictions through the variables  $W_i$  for technical reasons 238presented in Section 1.3 of SI. BSMs allowed us estimating the correlations between the errors of 239the land-use choices and ecological equations and therefore inferring the potential interactions 240between land-use choices and responses of tree species (Section 1.4 of SI). More details regarding 241the technical aspects of BSM predictions are provided in Section 1.5 of SI.
- BSMs are compatible with the generalized linear model (GLM) and with the generalized

243additive model (GAM) frameworks, as they both allow estimating  $f_i(X_i,W_i)$  jointly with 244  $f_p(X_i)$  and taking into account the potential dependence between equation errors. BSMs were 245therefore estimated in both GLM and GAM frameworks, using probit-linked functions. The GAMs 246analyses were performed to ensure the robustness of the GLM results, as GAMs reduce errors due 247to model misspecification and therefore reduce the risk of misleading interpretations of the error 248correlation between the ecological and economical equations (Ahn & Powell 1993). The BSMs 249estimated using GLMs and those estimated using semiparametric GAMs yielded quantitatively 250similar results (Section 3 of SI). For the sake of conciseness only the GLM results are presented in 251the main text. Note that selection biases are not restricted to the GLM or GAM cases but apply to 252any model assuming errors independently distributed from land-use choice.

## 253**2.4. Data**

# 2542.4.1 Land-use and species distribution data

Land-use and species distribution data were derived from the French national forest 256inventory which provides a systematic record of tree species presence/absence on a regular 1 km 257grid over the mainland territory (Figure 1 of SI) for sessile oak (*Quercus petraea*), pubescent oak 258(*Quercus pubscens*), common beech (*Fagus sylvatica*) and silver fir (*Abies alba*). This dataset 259therefore allowed separating land-uses in two categories: forest and non-forest. To test for the effect 260of spatial resolution on BSMs calibration and predictions, the dataset was upscaled on squared grid 261at three different resolutions 2 km, 4 km and 8 km sides, in accordance with the environmental data. 262The upscaling procedure was straightforward: a pixel indicated the presence (of a species or of the 263forested land-use) at the coarser resolution if it included at least one pixel where the species or the 264land-use was indicated present at the finer resolution.

## 2652.4.2 Environmental data

All the environmental variables were scaled at the different resolutions: 4 km<sup>2</sup>, 16 km<sup>2</sup> and 8

267km². Climate variables were derived from the SAFRAN re-analysis which includes temperature, 268rainfall, and radiation on a 3 hourly basis at 8 km² resolutions (Vidal et al 2010). These variables 269were averaged at a monthly time step and downscaled at 2 km² and 4 km² resolutions using a thin 270plate spline interpolation procedure with 3 predictors (elevation, latitude and longitude), 271implemented in the packages "fields" and "raster" in R (R Development Core Team 2012). This 272methodology has been validated with surface observations of temperature and rainfall over a region 273of southern France by Ruffault et al (2014). From the downscaled climatic variables, we derived 6 274variables considered critical to plant physiological function and survival, as in Cheaib et al (2012), 275which are summarized in Table 2 of SI. The slope and exposure data were computed by applying 276the "terrain" function of the "raster" package of R to digital elevation models at each resolution. 277The 1 km French soil data base developed by the INRA (Infosol Unit, INRA, Orleans, Jamagne et 278al. 1995) was averaged at the different resolutions and used to calculate soil water holding capacity 279(Cheaib et al 2012). To include realistic environmental conditions and reduce multicollinearity, we 280selected the first two axes of a principal component analysis (PCA) based on monthly climate 281variables and the first axis of a PCA made on soil and topographic variables (see Figure 2 of SI).

#### 2822.4.3 Economic data

Environmental predictors are also used in the econometric equation of land-use choices, in 284addition to some proxies of economic returns from the work of Ay *et al.* (2014). These authors 285approximated the monetary returns from crops by the land prices from the French ministry of 286agriculture in 2005, available at a regional scale named *Petites Régions Agricoles*. Monetary returns 287from forests were approximated by multiplying raw productions and unitary wood prices, divided 288by forest acreages. Monetary returns from urban area were approximated by population densities. A 289full description of the sample and the variables is reported in Table 1 and 2 of SI. Accessing 290monetary returns from different land uses is nevertheless not a necessary condition to estimate a 291BSM. Some other proxy variables could provide sufficient exclusion restrictions in numerous

292situations (such as FAO suitability index or NASA human footprint map).

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# 2943. Results

## 295**3.1.** Estimated parameters and response curves

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Explanatory variables included in the BSM all explained significantly the land-use choices 298and the tree species presences (Table 3 to 8 of SI). The estimation of the econometric equation (2) 299yielded adjusted R² ranging from 0.09 to 0.16 with economically intuitive effects: the probability of 300having a compatible land-use (i.e., forest) increased with the return from forests and decreased with 301the returns from cropland and urban areas. The estimation of the ecological equation (1) yielded 302adjusted R² ranging from 0.05 to 0.34 with significant bioclimatic predictors. We found a 95% 303significant selection bias for the four tree species at 2 and 4 km resolutions while at 8 km only 304*F.sylvatica* presents a significant selection bias. At 2 km resolution, the two Oak species present 305positive correlations of the errors (respectively 0.95 and 0.96 for *Q. petraea* and *Q. pubescens*, 306Table 3 of SI) while the two other species present negative covariance (respectively -0.75 and -0.82 307for *F. sylvatica* and *A. alba*, Table 3 of SI).

We simulated the effects of environmental predictors on the probabilities of species presence 309(response curves). Environmental factors were first projected in PCAs, and the first PCA axes were 310used as covariates (Figure 2 of SI). The predicted responses to a given covariate were computed by 311predicting the probability of presence while holding the other covariates at their sample means. The 312response curves estimated at 2 km resolution strongly differed between the BSM and the classical 313P-O and P-A SDMs for most of the species (Figure 1). The direction and magnitude of the deviation 314from the BSM response curves in the four species varied according to i) the sign of the correlation 315between the errors obtained from the environmental and economic equations and ii) the type of 316dataset used for the calibration of the classical SDMs (P-O or P-A). On the one hand, the two oak

317species, with positive correlation presented a similar pattern: the classical P-A SDMs overestimated 318the probabilities compared to the BSM whereas classical P-O SDMs yielded response curves very 319close to those of the BSM (Figure 1, panel A and B). On the other hand, for the two other species 320with negative correlation the response curves obtained with either the P-O or the P-A SDMs 321strongly underestimated the probabilities of presence compared to the BSM predictions (Figure 1, 322panel C and D). For these latter species, the magnitude of the deviation from the BSM was always 323greater for P-O SDMS.

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# 325[Expected location of Figure 1]

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# 3273.2. Prediction errors and spatial distribution

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We compared the probabilities predicted by the BSM and by the classical P-O and P-A 330SDMs, at the 2 km, 4 km and 8 km resolutions. We used the root mean of squared error (RMSE) to 331quantify the differences in predictions (eqn 12), taking the BSM predictions at 2 km as reference:

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$$RMSE_{m,r} = \sqrt{\frac{\sum_{i=1}^{n} (Prob_{i,m,r} - Prob_{i,BSM,2km})^{2}}{n}}$$
 (12)

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Where Prob is the estimated probability of presence, m={P-O SDM, P-A SDM, BSM}, r={2 336km, 4 km, 8 km resolution}, n is the number of sites. For a given resolution, the estimates of P-O 337and P-A SDMs strongly diverged from the estimates of BSM for most species, yielding high 338RMSEs (Table 1). Decreasing the spatial resolution had very strong effects on the prediction of all 339SDMs, and on the resulting RMSE. The effect of the spatial resolution on SDM predictions is

340indeed of the same order of magnitude than the effect related to the choice of the modelling 341approach (BSM or classical SDMs).

342

# 343 [Expected location of Table 1]

344As an external source of data about tree species potential presence, we use the EuroVegMap dataset 345(www.floraweb.de/vegetation/dnld\_eurovegmap.html) to provide an additional evaluation. Such 346data could be considered closer to the potential distribution relatively to IFN data because they were 347built at larger scale, upon an ensemble of criteria (e.g. presence of accompanying species, soil type 348and climate type) that may discriminate if an area of land is suitable for a given species 349independently from current land use. Figure 18 in SI 3.9 shows that, for species with negative 350correlations (*F.sylvatica*, *A.alba*), the relative performance of BSM increases highly, and could even 351slightly exceeded the levels of classical SDM for *A.alba*. This shows that BSM perform better at 352explaining the differences between the two data sources than classical SDMs.

353

The spatial patterns of predicted potential distribution differed markedly among the different 355models for each species at 2 km (Figure 2). This resulted in a bias (see Figure 15b in SI 3.8) 356affecting the predicted species distributions over the North Eastern-part and the Southern part of the 357territory in *Q. petraea* and *Q. pubescens*, respectively. Classical SDMs appeared to strongly 358underestimate the potential presence of *F. sylvatica* and *A. alba* with the exception of mountain 359areas where much smaller distribution areas were predicted.

360

## 361[Expected location of Figure 2]

362

## 3633.3 Interaction between ecological and economical gradients

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The additional information contained in BSM compared to classical SDMs can be assessed 366by computing the covariance between the ecological gradient  $\mu_i$  and the economic gradient  $\tilde{u}_i$  . 367The total correlation  $\rho_T$  between the variables describing potential tree species presence and 368compatible land-use is decomposed between an observed part ( $\rho_O$ , due to the included 369predictors) and an unobserved part ( $\rho_U$ , due to errors' correlation) as formally shown in section 3701.4 of SI:

371 
$$\rho_{T} = \frac{cov(\mu_{i}, \tilde{u}_{i})}{\sigma_{T}} = \frac{cov(f_{p}(X_{i}), f_{l}(X_{i}, W_{i}))}{\sigma_{T}} + \frac{cov(\varepsilon_{i}, \xi_{i})}{\sigma_{T}} = \rho_{O} + \rho_{U} \quad (13)$$

A positive  $\rho_O$  indicates that the predictors have the same effects on ecological suitability 373and economic incentive of having a forested land-use, and this was the case for all species and for 374all the resolutions (Table 2). By contrast, the sign of  $\rho_U$  varied between tree species. An 375important result is that  $\rho_U$  had a greater influence on the total correlation than  $\rho_O$ : for every 376species,  $\rho_T$  and  $\rho_U$  had the same sign. This implies that neglecting the correlation between 377errors (as it is typically done by classical SDMs) induces a strong misestimation of the interactions 378between the ecological and economical gradients.

379

# 380[Expected location of Table 2]

381

16

In the case of species with positive total correlations (*i.e. Quercus*), sites with high 383probability of presence are also sites with high probability of compatible land-use (Figure 3). This 384indicates a synergy between the two dimensions: the best suitable sites for the species are actually 385forested. The opposite is obtained for species with negative correlation, for which the hidden part of

386the niche is composed of sites with above-than-average probabilities of potential presences (Figure 3873). This indicates an antagonism between the ecological and the economical dimensions. Similar 388results were found at 4 km and 8 km resolution, highlighting the importance of  $\rho_T$  to 389characterise the interaction between ecological and economical gradients and the importance of 390BSMs to estimate it.

391

## 392[Expected location of Figure 3]

393

## 3944. Discussion

This study examines the influence of human decisions about land-use on the calibration and 396prediction of tree species distribution models (SDMs). We did so by developing a bivariate selection 397model (BSM) that is based on the coupling of an ecological equation of species distribution and an 398econometric equation of land-use choices. An empirical application on four tree species using 399standards statistical methods (GLMs and GAMs) illustrates and confirms the theoretical 400developments. It shows that classical SDMs may, depending on the species and the spatial 401resolution of the calibration dataset, either over or underestimate the potential probabilities of 402presence. We show that the bias is important at fine spatial resolution and decreases at coarse 403resolution. Hereafter, we discuss the causes of between-species variations and spatial resolution on 404the direction and magnitude of the bias. Finally, we discuss the implications of coupling ecological 405models of species distributions with econometric models of land-use choices.

406

# 4074.1 Land-use selection bias affects the predictions of potential presence

We found a significant spatial correlation between the economic requirements of the forested 409land-use and the ecological requirements of every studied tree species. It is long recognized that in

410anthropogenized areas, forests are preferentially located on poor soils with low water holding 411capacity, where no profitable agriculture is possible (Ricardo 1817). Therefore, tree species tolerant 412to such constraining environmental conditions will be more frequently found at forested sites than 413species with ecological requirements that match conditions favouring high economic returns for a 414different land-use (e.g. agriculture). This is well illustrated in our results: for *Quercus* species -- 415which are characterised by a relatively high tolerance to dry sites (Arend *et al.* 2011; Michelot *et al.* 4162012; Lévesque *et al.* 2014, see also Figure 2) -- we found a positive correlations between the errors 417of the ecological and economical equations (Table 3). By contrast, for *Abies and Fagus* we found a 418negative correlation between errors of the two equations (Table 2). This is consistent with the fact 419that these latter species are relatively more sensitive to dry climate and shallow soil (Aranda et al 4202005), and grow preferentially on fertile sites of temperate areas which are therefore predominantly 421used for agriculture.

In agreement with the theory developed in this study, the correlation between ecological and 423economical gradients found in every species resulted in a significant land-use selection bias that 424affected the predictions of classical SDMs. We found that P-O SDMs always underestimate the 425probability of presence, whatever the species considered (Figure 3). All sites with land-uses 426incompatible with forests are indeed considered as absences in P-O SDM calibration, which 427consequently keeps some of the environmental conditions that are suitable for a given species out of 428the predicted potential niche. Contrastingly, the sign of the bias affecting P-A SDM predictions 429varies among species: positive bias in the two *Quercus* species and negative bias in *Abies* and 430*Fagus* (Table 2). Unlike P-O SDMs, P-A SDMs characterize the environmental conditions of 431unsuitable sites in order to predict species distributions and niches (Lobo *et al.*, 2010). However, if 432the ecological requirements of a tree species are preferably found at locations where land-use is 433incompatible (i.e., there is a negative correlation between ecological and economical gradients), the 434proportion of observed absence data is higher than in the case of a randomly distributed species. In

435such case, P-A SDMs underestimate the predicted probability of presence (Figure 3, *Abies* and 436*Fagus*). Conversely, if the ecological requirements of a tree species are preferably found at locations 437where land-use is compatible, P-A SDMs overestimate the predicted probability of presence (Figure 4383, *Quercus* sp.).

Overall, our results demonstrate that the land-use selection bias can lead to strong spatial 440miss-estimations of the predicted probabilities of presence of different tree species (Figure 2). Not 441accounting for this bias may have great implications for the many studies that deal with the 442probability of presence derived from classical P-O-SDM or P-A-SDM dataset in anthropized area. 443Furthermore, it is worth mentioning that such a bias will affect the predicted distribution of any 444species that is more specifically found within a certain land-use. Consequently, the projections of 445future distribution areas, that are a major tool for conservation strategy (Wilson *et al.* 2011), are 446likely to be biased as well. It is crucial to consider land-use selection bias in SDMs if we are to 447provide sound prediction of the future of species distributions in Western Europe and other 448anthropized areas.

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## 450**4.2.** How to evaluate SDMs in anthropized areas?

Our results question the relevance of the validation of statistical models based on presence-452absence data from forest inventories. Indeed, the maps of probabilities computed with the BSM 453significantly differ from the presence-absence (P-A) records used to calibrate them, and such 454differences are far more important than for both classical SDMs (Figure 2, section 3.9 of SI). We 455argue that by accounting for the land-use selection bias, our BSM yields probabilities more closely 456related to the unobserved potential niche of the species than a classical SDM. This interpretation is 457consistent with (i) studies proposing that part of the differences between the realized and the 458potential niche is shaped by land cover modification related to human activities (Svenning et al

4592010); and ii) with studies that found a significant role of land-use in the distribution of plant 460species independently of climate soil or topographic variables (e.g. Duckworth et al 2000).

It would be of great value to find new ways of validation/assessment for distribution models 461 462based on independent data. Such an evaluation is nevertheless quite a hard task given the high 463discrepancies reported when multiple sources of tree distribution data are compared (Duputie et al., 4642013). Evaluation over pre-historical periods, when vegetation cover was not affected by human 465induces modifications of the land covers, by using distributions databases built from paleo materials 466may be a solution (Svenning et al., 2011). However paleo-data are subjected to uncertainty far more 467important than classical inventory data (Svenning et al., 2011). Another interesting way forward is 468the use of process-based models, including dynamic global vegetation models, that capture the 469physiological mechanisms determining plant survival and reproduction (Dormann et al. 2012). 470Because these models are based on the biophysical rules driving plant functioning, we expect their 471predictions to be unaffected by land-use selection bias, when they are calibrated on experimental 472data (Benito Garzón et al. 2011; Anderegg et al. 2015). In such case, they provide valuable estimate 473of the potential niche of the species (Morin et al. 2007; Hickler et al. 2012). A recent study 474conducted over France (Cheaib et al. 2012), showed that species distribution estimates from 475process-based models are close to the distribution observed in temperate oaks, but predict a larger 476distributional area than what is observed in common Beech. These additional presence areas 477correspond to regions that are mostly covered by agricultural land-use (e.g. Beauce region in 478Southern Paris, Normandie in North West France). This qualitative assessment echoes the results 479obtained in our theoretical development about the sign of the correlation between errors. Indeed, 480Quercus exhibits a positive correlation sign and is thus weakly affected by land use selection bias, 481whereas *Fagus* shows highly negative correlation leading to an important selection bias.

However, we should recall that process-based model predictions remain contingent on the 483hypothetical biological processes considered, and that they may be more uncertain than correlative

484approaches estimates (Buckley *et al.* 2010). Promising attempts have been made to circumvent the 485limitations of individual modelling approaches, either in building hybrid models with both 486statistical and mechanistical features (Dormann *et al.* 2012), or by combining the predictive power 487of several independent models (Romero *et al.* 2015). These advances will likely help reducing the 488bias currently affecting SDM predictions. Given the current state of the art in species distribution 489modelling, comparing the simulations of a variety of approaches (Gritti *et al.* 2013) against multiple 490sources of data -- including expert knowledge --remains necessary to ensure adequate evaluations of 491model outputs (Cheaib *et al.*, 2012; Duputie *et al.*, 2013).

# 4.3 The importance of land-use selection bias increases at fine resolution

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493 An increasing number of studies have pointed out the importance of working at fine spatial 494resolution to properly assess the outcome of climate change on species persistence (Trivedi et al 4952008; Randin et al 2009). These scientific results are followed by an increasing demand from 496applied researchers for fine resolution data. This leads to a growing importance of the land-use 497selection bias presented in this study, as we showed an increasing importance of the land-use 498selection bias with the spatial resolution (Table 2). Our results are in agreement with other studies 499showing that land-use can help to improve the explanatory power of species distribution models 500based on climate variable, mostly at fine resolution (Pearson et al 2004; Luoto et al 2007). More 501generally our results are in line with the premise that different factors shape the distribution of 502species according to a hierarchy of spatial scale; in which climate is more important at coarse scale 503and land-use at finer scale (Pearson et al 2003). However and more importantly, the changes in the 504shape of the response curves to environmental variables between fine and coarse resolution have 505likely great implications for climate projections (Figure 2). Indeed we can reasonably expect that 506the projections under climate change scenario based on BSM calibrated at fine resolution will 507deeply differ from the projections based either on classical SDMs at fine resolution (without land-508use) or on BSM at coarse resolution. Such pronounced effect of land-use contrasts with the different

509studies showing little changes in the projected probabilities when land-use is accounted for (Barbet-510Massin et al 2012b; Martin et al 2013). Hence, our work highlights the need to jointly estimate 511land-use choices and species responses to bioclimatic variables in a bivariate context to capture 512their interacting effects on species distribution (Clavero *et al.* 2011; Mantyka-pringle *et al.* 2012) 513and to deal with the land-use selection bias affecting the SDM calibration in human-dominated 514ecosystems. Consider for instance *F. sylvatica*, a major European species whose distribution is 515predicted to strongly shrink due to climate changes (Cheaib *et al.* 2012). It is in fact probable that 516some suitable areas will last on the territory for decades, but we highlight in this study that the best 517sites for this species are often not under a compatible land-use (i.e. they are not forested areas). So 518maintaining *F. sylvatica* in France will have a cost that it is necessary to anticipate in order to refine 519the predictions of the outcomes of global changes on species distributions (Fisher *et al.* 2011).

520

## 521**4.4** Going further the selection bias: toward an integrated projecting

The BSM that we present here constitutes an important step to account for the economic forces 523shaping the observed species distributions. Selection bias is proven to be important for tree 524distribution in France (Table 1) but other types of bias could persist in modelling species 525distributions. One typical example is the high proportion of absence for species with a highly 526negative correlation. There are some alternative models based on zero-inflated probability 527distributions that could be used and even mixed with the BSM to correct for this. We believe that 528this issue would deserve proper researches to improve modeling practices.

Our empirical estimates of the interactions between economic and ecological gradients (Figure 5303; Table 2) revealed how the studied species are differently influenced by environmental and 531economic variables. The use of explicit economic data to predict land-use paves the way for more 532direct quantification of synergies and trade-off between environmental protection and economical 533welfare. Using BSM for future projections imply a qualitative change of the role of economic 534variables. From their technical role in removing the selection bias, economic variables become key

535predictors for future land use choices. Hence, instead of using exogenous global/regional scenario 536of future land-use (as generally done is SDM, e.g., Martin et al., 2013), the BSM would be based on 537the trajectory of monetary returns from different land-use and may help to quantify the cost for 538conservation of a given species. In our opinion, this alternative approach is promising to help 539decision making by policies.

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# **Supporting Information:**

- **1. Theory of the Bivariate Selection Model** 
  - 2. Data description
- **3. Additional results**

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544

547**Biosketch :** The authors belong to a multidisciplinary project founded by the French government 548(LabEx BASC) that aims to provide new knowledge regarding species distribution drivers and to 549design innovative guidelines toward sustainable land management. The present team included both 550ecologists interested in the functioning of plants in relation to the environment (Paul Leadley, 551Nicolas K. Martin-StPaul and Joannès Guillemot) and economists specialized in land-use 552economics and its influence on biodiversity (Luc Doyen, Jean-Sauveur Ay). The authors share an 553interest in the understanding of species distribution in anthropized areas, and in the use of SDM to 554gain knowledge and inform environmental decisions.

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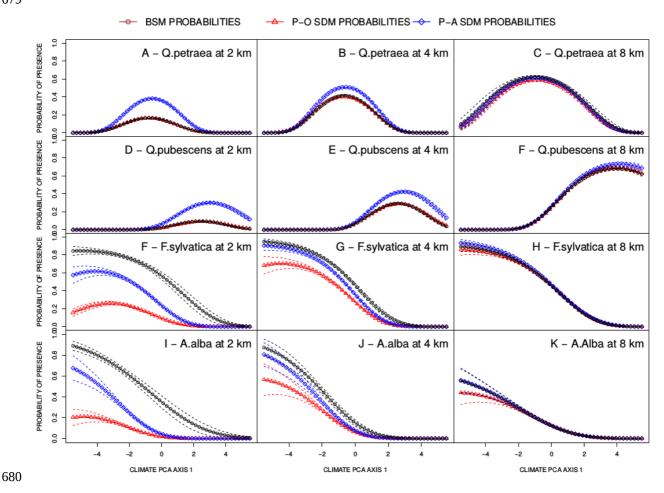
**Table 1:** Comparison of the probabilities predicted by the three modelling approaches: BSM, (P-O) 667and (P-A) SDMs. Figures are RMSEs calculated using BSM predictions at 2 km as reference.

Species	Scale	BSM	РО	PA
Q.petraea	$2 \mathrm{km}$	0.00	0.00	0.19
	$4 \mathrm{km}$	0.17	0.17	0.28
	$8\mathrm{km}$	0.46	0.39	0.43
Q.pubescens	$2 \mathrm{km}$	0.00	0.10	0.23
	$4 \mathrm{km}$	0.14	0.14	0.19
	$8 \mathrm{km}$	0.29	0.29	0.31
F. sylvatica	2km	0.00	0.48	0.35
	$4 \mathrm{km}$	0.24	0.41	0.34
	$8\mathrm{km}$	0.33	0.33	0.31
A. alba	2km	0.00	0.33	0.29
	$4 \mathrm{km}$	0.28	0.32	0.32
	$8\mathrm{km}$	0.31	0.30	0.31

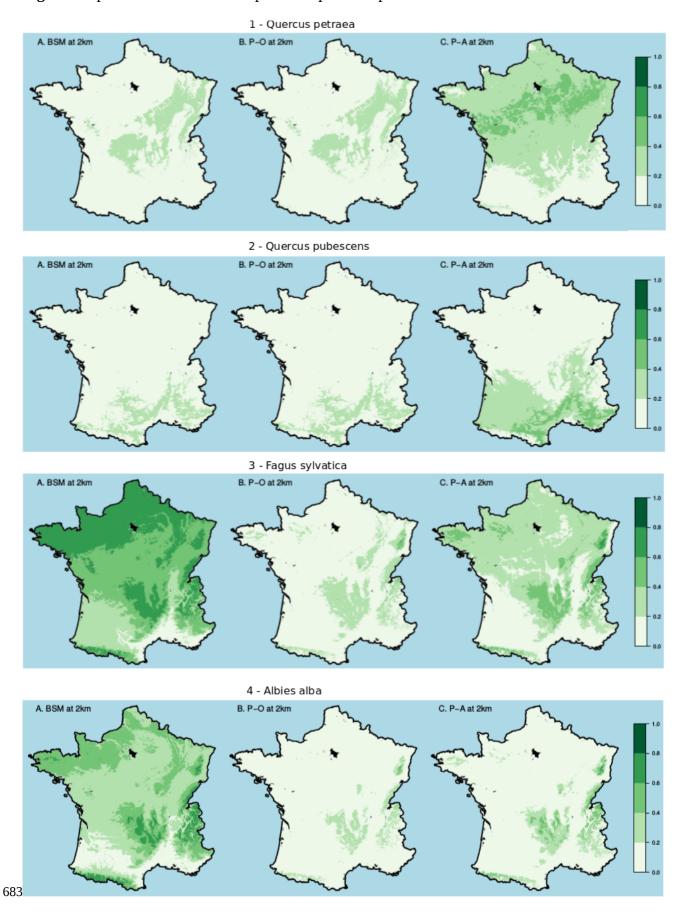
**Table 2**: Decomposition of the spatial covariance between ecological and economical 669gradients.  $\rho_T$  is the total correlation between ecological and economical components. It 670is decomposed into an observed part  $\rho_0$  (due to the included predictors) and a 671unobserved part  $\rho_0$  (due to the errors' correlation). The terms in brackets represent the 672confidence intervals at 95%.

		TOTAL		OBSERVED		UNOBSERVED	
Resol.	Species	$ ho_T$	IC 95%	$\rho_O$	IC 95%	$ ho_U$	IC $95\%$
2 KM	Q.petraea Q.pubescens F.sylvatica A.alba	0.707 $0.781$ $-0.568$ $-0.620$		0.143 0.180 0.007 0.041	$ \begin{bmatrix} 0.06, 0.23 \\ [0.1, 0.27] \\ [-0.04, 0.05] \\ [-0.09, 0.17] \end{bmatrix} $	0.565 $0.600$ $-0.575$ $-0.660$	$ \begin{array}{c} [0.53, 0.58] \\ [0.55, 0.62] \\ [-0.62, -0.51] \\ [-0.7, -0.61] \end{array} $
4 KM	Q.petraea Q.pubescens F.sylvatica A.alba	0.567 $0.727$ $-0.325$ $-0.246$	$ \begin{bmatrix} 0.35, 0.72 \\ [0.52, 0.87] \\ [-0.43, -0.21] \\ [-0.58, 0.11] \end{bmatrix} $	0.136 0.226 0.078 0.153	$ \begin{bmatrix} 0.04, 0.23 \\ [0.1, 0.35] \\ [0.03, 0.12] \\ [-0.1, 0.41] \end{bmatrix} $	0.431 $0.502$ $-0.403$ $-0.399$	
8 KM	Q.petraea Q.pubescens F.sylvatica A.alba	-0.307 $0.712$ $0.520$ $0.258$	$   \begin{bmatrix}     -0.56, 0.14 \\     -0.43, 0.75 \end{bmatrix} \\   \begin{bmatrix}     0.21, 0.75 \\     -0.11, 0.63 \end{bmatrix} $	0.020 0.163 0.153 0.213	$   \begin{bmatrix}     -0.03, 0.07 \\     0.12, 0.2 \\     0.02, 0.29 \\     0.01, 0.42   \end{bmatrix} $	-0.327 0.549 0.366 0.044	$   \begin{bmatrix}     -0.53, 0.07 \\     -0.55, 0.55 \end{bmatrix} \\   \begin{bmatrix}     0.19, 0.47 \\     -0.12, 0.21 \end{bmatrix} $

**Figure 1.** Predicted response curves of the presence probabilities to the environmental gradient 678(first axis of the PCA on environmental variables, see Figure 2 of SI). 679



**Figure 2**. Spatial distribution of the predicted potential presence at 2 km resolution.



**Figure 3:** Probabilities of potential presence of tree species according to the presence of compatible 685land use (i.e., forested land use). 686

