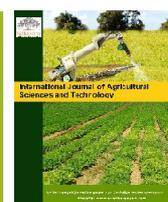




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Ecological niche modeling to assessment of potential distribution of *Neodiprion abietis* (Harris, 1841) (Insecta, Hymenoptera, Diprionidae) in Eurasia

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Abstract

In the article first assesses the potential distribution in Eurasia of *Neodiprion abietis* (Harris, 1841) first time assessed. The species is widely distributed in North America fir and spruce defoliator, intercepted in 2016 in the Netherlands. Analysis of the literature data on the known distribution and host plants of the species is given. Insufficient data for ecological modeling is shown, which causes a high degree of uncertainty in assessment of potential range. The results of modeling the potential range of the species using several algorithms are presented. The reliability of the obtained models and the importance of bio-climatic environmental variables for species distribution are assessed.

Keywords: Niche modeling, Potential distribution, Forest pests, Conifers, *Neodiprion abietis*

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1. Introduction

Neodiprion abietis (Harris, 1841) (Insecta, Hymenoptera, Diprionidae) is a serious pest of coniferous trees, widely distributed in North America from Canada to Mexico (GBIF 2020a), *Neodiprion abietis* (NEODAB) [World distribution]. The larvae of this sawfly cause defoliation (sometimes severe) of the host plants, which cause their weakening, decreasing growth, and sometimes tree death (Piene *et al.*, 2001). The species has several ecological and geographical forms feeding on various fir and spruce species, including European spruce (*Picea abies* (L.) H.Karst.) (Knerer and Atwood, 1972). In autumn 2016, the cocoon *N. abietis* was detected in the Netherlands on branches of *Gaultheria sp.* imported from the USA. (Interception..., 2017). For this reason, in 2017 the species was included in the Alert list of the European and Mediterranean Plant Protection Organization. To assess the possibility of *N. abietis* acclimatization in the Old World, an attempt was made for the first time in this article to predict its potential distribution by the ecological niche modeling based on several algorithms.

2. Materials and methods

The source of data on the occurrences of *N. abietis* in North America were GBIF dataset (GBIF, 2020b). The distribution of potential host plants was reconstructed from various sources mentioned in the relevant section of the article. The free software openModeller desktop (de Souza Muñoz *et al.*, 2011) version 1.1.0 and Maxent (Steven *et al.*, 2020) version

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3.4.1 were used for ecological niche modeling of the potential *N. abietis* distribution. Data on bioclimatic environmental variables were obtained from raster layers of WorldClim v. 2.1 representing average values of climate data for 1970-2000 (Fick and Hijmans, 2017). Data processing and mapping were performed in QGIS (QGIS.org), statistical analysis—in Microsoft Excel. Layers of administrative boundaries are based on GADM database (www.gadm.org) version 3.4. The results of the ecological niche modeling experiments and the original shape-files on which the figures for this article are based are available in the attached dataset.

3. Results and discussion

There is no doubt that the main factor limiting the distribution of phytophagous is the presence of host plants. Unfortunately, their exact list for *N. abietis* has not yet been identified. Various authors mentioned plants of the genus *Abies*, *Picea*, *Tsuga*, *Pseudotsuga*, *Pinus* and even *Thuja* as such (Bird, 1929). Ross (1955) identified the *abietis* complex in the *sertifer* group of the *Neodiprion* genus, including two species, one of which inhabits *Abies* and *Picea*, the other (not formally described) - *Tsuga* and *Pseudotsuga*. Knerer and Atwood in their description of biology “*Neodiprion abietis* Complex” (Knerer and Atwood, 1972) identified four forms (“strains”), among the host plants mentioned balsam fir (*Abies balsamea* (L.)), white fir (*Abies concolor* (Gordon) Lindley ex Hildebrand), white spruce (*Picea glauca* (Moench) Voss), black spruce (*Picea mariana* (Mill.) Britton, Sterns & Poggenburg), *European spruce* (*Picea abies* (L.) H. Karst.). However, it cannot be said that these ecological and geographical diversity of the species is limited to these four forms. Moreover, the identity and limits of the *N. abietis* species need to be clarified, and all reports on findings of this species published in the 19th and early 20th centuries (before the description of closely related species and the revision of the genus prepared by Ross) should be considered critically. Due to the high uncertainty of the list of potential host plants of *N. abietis*, all species of the genus *Abies* and *Picea* are considered here as such. Reports on nutrition by plants of other genera are questionable, and most likely a mistake.

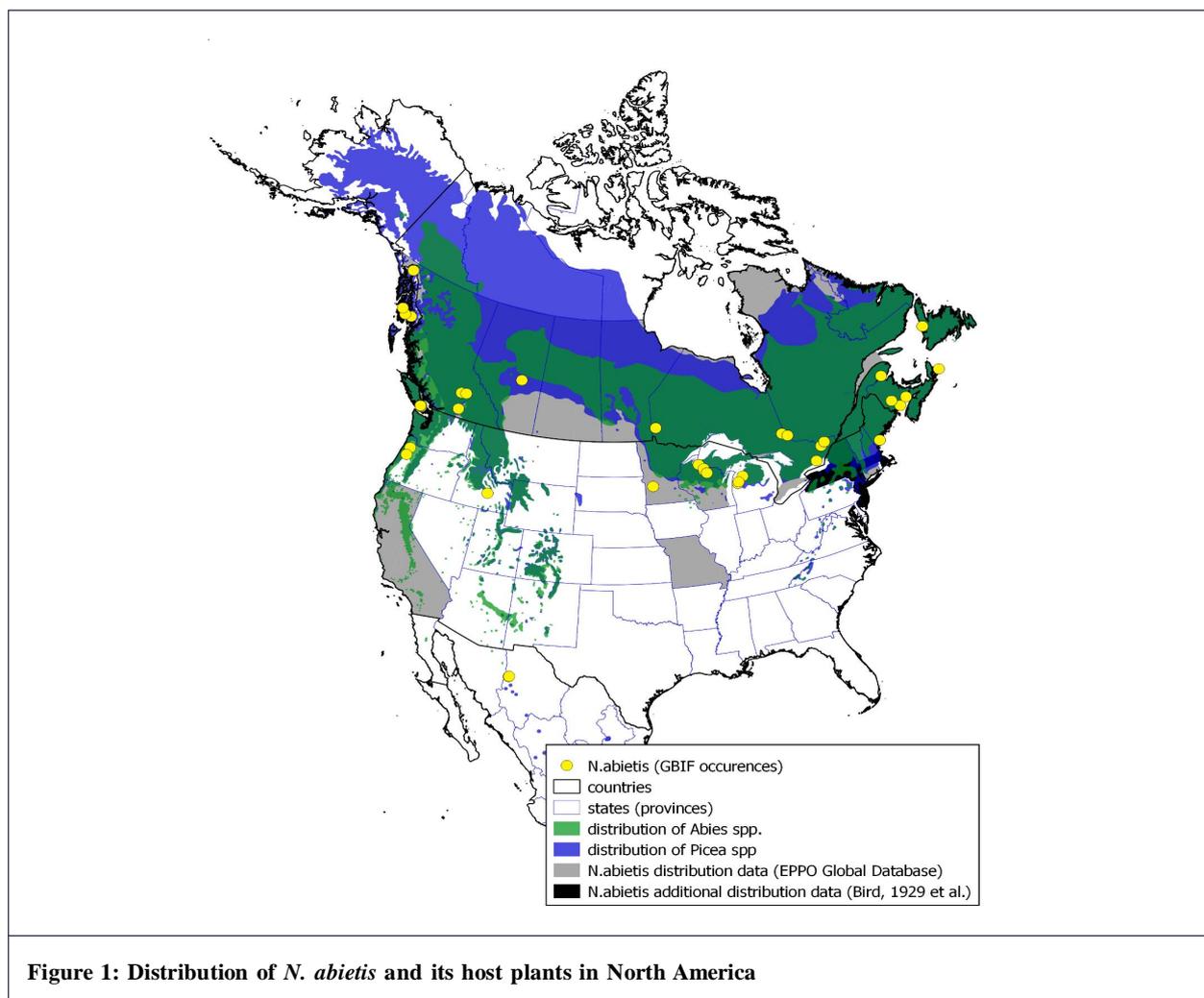
Data on host plant distribution in North America were obtained from clone of digital representations of tree species range maps from “Atlas of United States Trees” by Elbert L. Little, Jr. (Petry, 2020). Fir and spruce ranges within Europe were established with some approximation from “Tree species maps for Europe” (Brus *et al.*, 2012) and “Chorological data for the main European woody species” (Caudullo *et al.*, 2020). The distribution of these genera in Asia was reconstructed on the data from the “Agricultural Ecological Atlas of Russia and Neighboring Countries” (Afonin *et al.*, 2008), maps of the USSR tree ranges (Sokolov *et al.*, 1977), some other publications (Xiang *et al.*, 2007; Lockwood *et al.*, 2013), and occurrences from *Abies* and *Picea* (GBIF 2020c; 2020d). Further assessment of potential *N. abietis* distribution in Eurasia was made within the limits of this approximate host plant range.

To assess the representativeness of the GBIF data on the occurrences of *N. abietis*, a map was made, correlating these occurrences with the literature data on the regions for which the species was reported, and the distribution of the host plants (Figure 1).

From this map it is clear that the GBIF dataset covers only part of the *N. abietis* range. For this reason, all models of potential distribution based on it are quite approximate and have a high degree of uncertainty. In further assessment of the reliability of various models, it was taken into account how adequately they describe the current distribution of the species, taking into account the literature data.

To select environmental variables optimal for niche modeling, the values of 19 bioclimatic variables from WorldClim raster layers were obtained for each of 121 points of occurrences from GBIF. The statistical analysis has shown correlation between these variables inside the dataset. For further modeling, 13 variables with Pearson correlation coefficient between them being less than 0.8 were selected:

- bio01 - Annual mean temperature;
- bio02 - Mean diurnal range (Mean of monthly (max temp - min temp));
- bio03 - Isothermality (BIO2/BIO7) (×100);
- bio05 - Max temperature of warmest month;
- bio07 - Temperature annual range (BIO5-BIO6);
- bio08 - Mean temperature of wettest quarter;
- bio09 - Mean temperature of driest quarter;
- bio10 - Mean temperature of warmest quarter
- bio12 - Annual precipitation;



bio13 - Precipitation of wettest month;

bio15 - Precipitation seasonality (coefficient of variation);

bio17 - Precipitation of driest quarter;

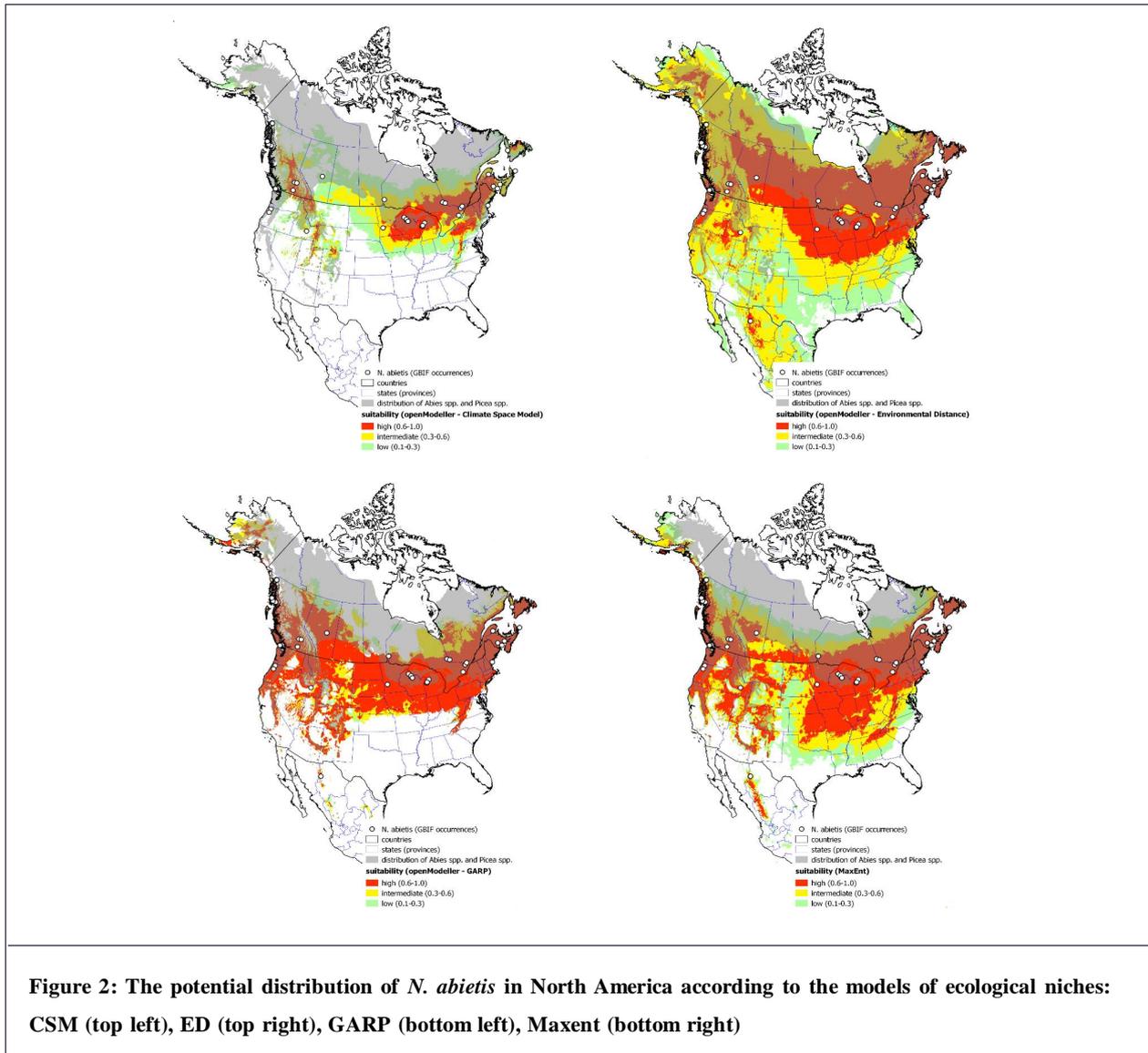
bio18 - Precipitation of coldest quarter.

Based on these variables, four models of the potential range were constructed using algorithms:

1. Climate Space Model (openModeller desktop, with default settings, AUC = 0.927) (CSM);
2. Environmental Distance (openModeller desktop, with default settings, AUC = 0.999) (ED);
3. GARP with best subsets - DesktopGARP implementation (openModeller desktop, with default settings, AUC = 0.987) (GARP);
4. Maximum Entropy (Maxent, settings were determined experimentally by the maximum AUC value (feature types used: hinge, product, linear, threshold, quadratic; maximum background points: 50000; maximum iterations: 5000; default prevalence: 0.9), AUC = 0.989) (Maxent).

The results of the experiments for North America are shown in Figure 2:

The CSM and ED models appear to be insufficiently reliable (despite the high AUC value). The CSM indicates an extremely limited potential range, not even covering the occurrences of the original dataset. The ED, on the contrary, shows an extremely wide potential distribution going far into polar latitudes. Thus, the GARP and Maxent models that describe the current distribution of the species more adequately are more reliable. The GARP model assumes a somewhat more distant northward distribution of *N. abietis*, while the Maxent model, on the contrary, allows a greater southward distribution of the species.



Extremely limited amount of georeferenced data on species distribution, and the doubtfulness of many references in the literature do not allow making a reasonable choice between these models. The high probability that the species will live in Missouri according to the Maxent model (where it is reported) can hardly be proof of greater reliability. The book (Craighead, 1950) referred to by EPPO Global Database provides only a general distribution of the species in eastern North America: “This species is widely distributed from Quebec to Manitoba in Canada, and in the United States from New England westward to Minnesota and Missouri”. *N. abietis* is also reported for Missouri by Ralf Bird (1929), but in the book to which he refers (Bureau of Entomology, 1908), there is only one mention of the species (on the page he indicated): “The fir sawfly (*Lophyrus abietis* Harris) defoliated much of the balsam fir on the islands and mainland along the coast of Maine.” - That is, the reference (quoted in later literature) is actually to the state of Maine, not to Missouri. In addition, there are no natural fir or spruce forests in Missouri (Petty, 2020). All of these facts raise great doubts about the presence of *N. abietis* in this state.

Thus, the GARP model may be more reliable, but this can only be confirmed with more detailed data on the *N. abietis* distribution. At the moment, it is reasonable to consider both this model and Maxent to assess the potential distribution of the species. Figures 3 and 4 shows the potential distribution of *N. abietis* in Eurasia according to both models, and taking into account the distribution of plants from the genera *Abies* and *Picea*.

The figures above show that the models more or less similarly determine the possible distribution of *N. abietis* in the Old World. Only relative degrees of area suitability differ significantly. The assessment of the probability of the species’ distribution in the territory of China also sharply differs: the GARP model assumes the possibility of wide distribution of *N. abietis* with high probability in many provinces, Maxent—almost everywhere mostly low probability of habitat, and other shape of the potential range.

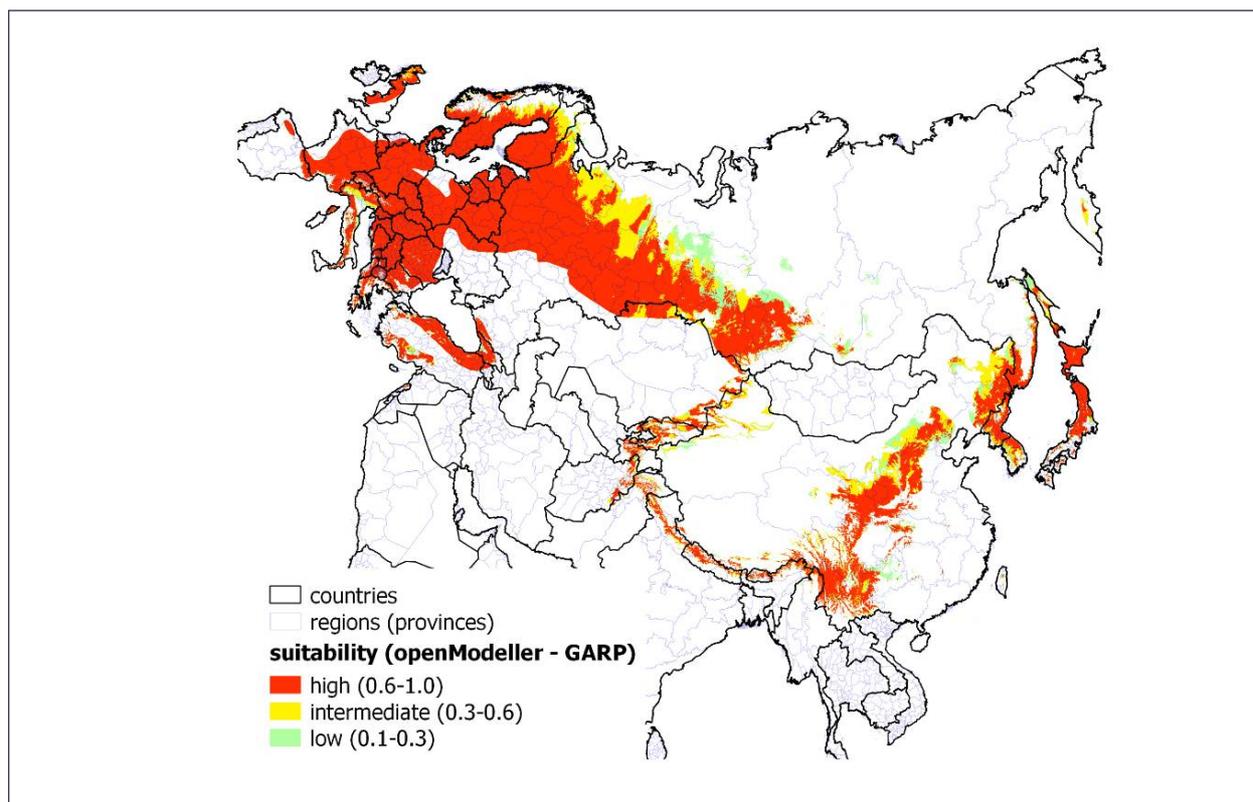


Figure 3: Potential distribution of *N. abietis* in Eurasia according to GARP model

Jackknife resampling in Maxent (Figure 4) showed the greatest importance for modeling the variables bio01 (annual mean temperature), bio10 (mean temperature of warmest quarter), bio17 (precipitation of driest quarter) and bio05 (maximum temperature of warmest month).

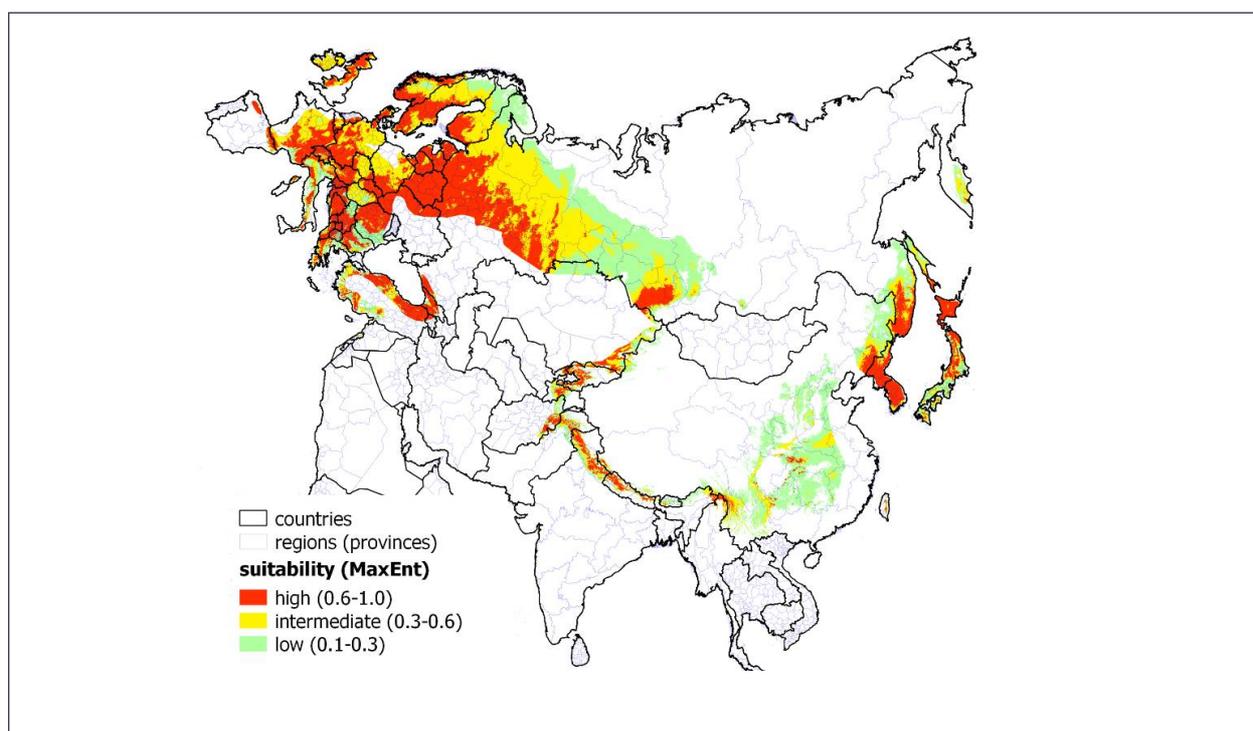


Figure 4: Potential distribution of *N. abietis* in Eurasia according to GARP Maxent model

This suggests that the limits of the ecological niche of *N. abietis* are determined (besides the presence of host plants) by relatively low annual and maximum temperatures without noticeable dry periods. This is more or less consistent with the modern distribution of the species shown here, as well as models of the potential distribution.

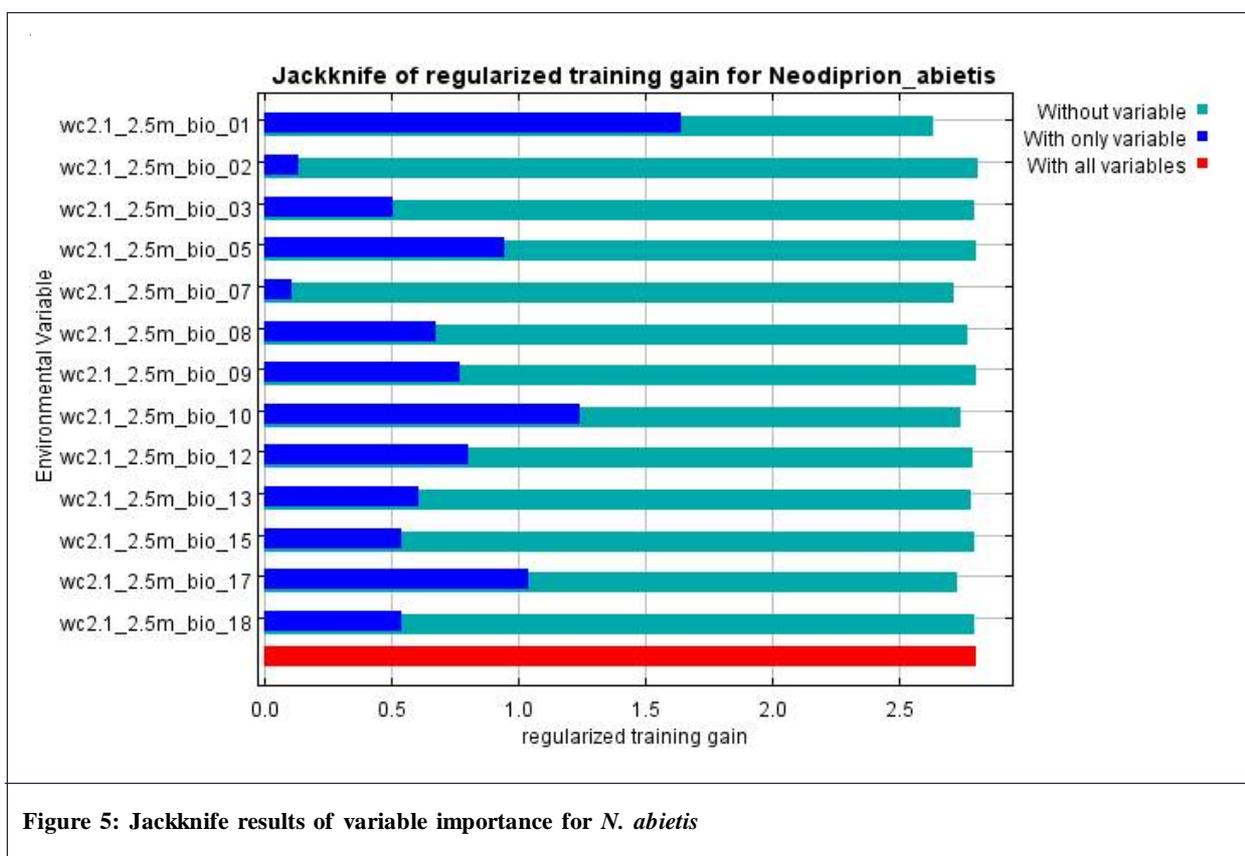


Figure 5: Jackknife results of variable importance for *N. abietis*

Conclusion

Modeling of the ecological niche of *N. abietis* showed a high probability of its distribution in Eurasia: most of Europe, Western Siberia, the Far East, Central Asia, the Himalayas, and possibly continental China. Due to the high uncertainty of both the list of host plants of the species and data on its current occurrence, the models shown for the potential distribution can only be considered as approximate. Indirect assessments suggest that models based on the genetic algorithm in openModeller and maximum entropy in Maxent are the most reliable. However, more accurate assessments are possible only on the basis of a more thorough study of the species in its natural range.

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