

Climatic extremes improve predictions of spatial patterns of tree species

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Understanding niche evolution, dynamics, and the response of species to climate change requires knowledge of the determinants of the environmental niche and species range limits. Mean values of climatic variables are often used in such analyses. In contrast, the increasing frequency of climate extremes suggests the importance of understanding their additional influence on range limits. Here, we assess how measures representing climate extremes (i.e., interannual variability in climate parameters) explain and predict spatial patterns of 11 tree species in Switzerland. We find clear, although comparably small, improvement (+20% in adjusted D^2 , +8% and +3% in cross-validated True Skill Statistic and area under the receiver operating characteristics curve values) in models that use measures of extremes in addition to means. The primary effect of including information on climate extremes is a correction of local overprediction and underprediction. Our results demonstrate that measures of climate extremes are important for understanding the climatic limits of tree species and assessing species niche characteristics. The inclusion of climate variability likely will improve models of species range limits under future conditions, where changes in mean climate and increased variability are expected.

climate change | ecological niche | generalized additive model | geographic range | species distribution models

The understanding of the principles and mechanisms that shape distribution patterns has long been a focus in biogeographical, ecological, and evolutionary research. The ecological niche concept, coined and initially developed by Grinnell (1), is the foundation for our understanding of the processes that shape the geographical distributions of species (2). Conceptual clarifications with regards to using the concept for the explanation of species ranges have been presented by several authors (3, 4). Climatic variables are often used to predict biogeographical patterns (5), and considerable effort has been put into improving methods to describe the response of species along climate gradients (6–8). These methods of species distribution or niche modeling are frequently used for conservation management (9–12), prediction of the likely effects of global change (13–16), and, increasingly, assessment of niche characteristics in the study of niche evolution (17–20). These studies in general use monthly or annual climatic means to analyze species distribution patterns. To date, little attention has been paid to the question of how climatic extremes, i.e., the long-term, interannual variation around mean values, could help to explain species distributions. There are two major reasons that highlight the importance of including climatic variability in niche analyses and models. First, ongoing climate change not only affects means but also extremes (21). Second, niche evolution often results in changes of the stress tolerance of evolving clades (22, 23). Thus, both adaptation and possible future response of species to climate are certainly affected by extremes in addition to means.

On a more fundamental level, the question is what aspects of climate actually determine the “climate response” of plant

species. The expected effects of climate extremes are diverse. At the range limit of species, source-sink dynamics likely exert strong influence (3, 24–26). In sessile organisms like plants, extremely adverse climate can constrain regeneration and impact range limits through increased mortality (27). Extremely favorable climate, in contrast, might allow unusually high reproductive success or the advance of range limits, especially when such effects occur over several years. The observed effects of severe heat and drought illustrate such range contractions at the trailing edge of species ranges. Such effects are visible by the dieback of Scots pine (*Pinus sylvestris* L.) in the warmest part of the dry valleys of the European Alps and increased diebacks in conifers in western North America (28) that are likely caused by recent warming and increased drought. Ongoing climate change impacts species phenology (29), individual growth (30), and ecosystem dynamics (31, 32). In addition, species have experienced recent range shifts (33–35), some of which are clearly induced by climatic extremes (36), whereas other responses seem more gradual (37). In particular, climatic extremes can be responsible for dieback at the trailing edge of species distributions (38). These examples illustrate the potential importance of including climatic variability into analyses and models.

Climatic variability alone likely does not explain the climatic response of species very well. Rather, variability may complement a species' response to mean climate, which summarizes complex mechanisms that are not directly expressed in a fitted response curve. For instance, under optimal temperature or moisture conditions, climatic variability or extremes could have relatively little effect on species, whereas the effect of variability is likely severe where mean temperature or water availability is closer to physiological tolerances. In contrast, under conditions in which warm temperatures and abundant moisture are favorable for growth, the effects of competition (on abundance, reproductive success, etc.) may outweigh the direct effects of climate. For example, a species' range might in some areas be directly limited by physiological tolerance to low temperatures, whereas in other areas with relatively warm climate, the range

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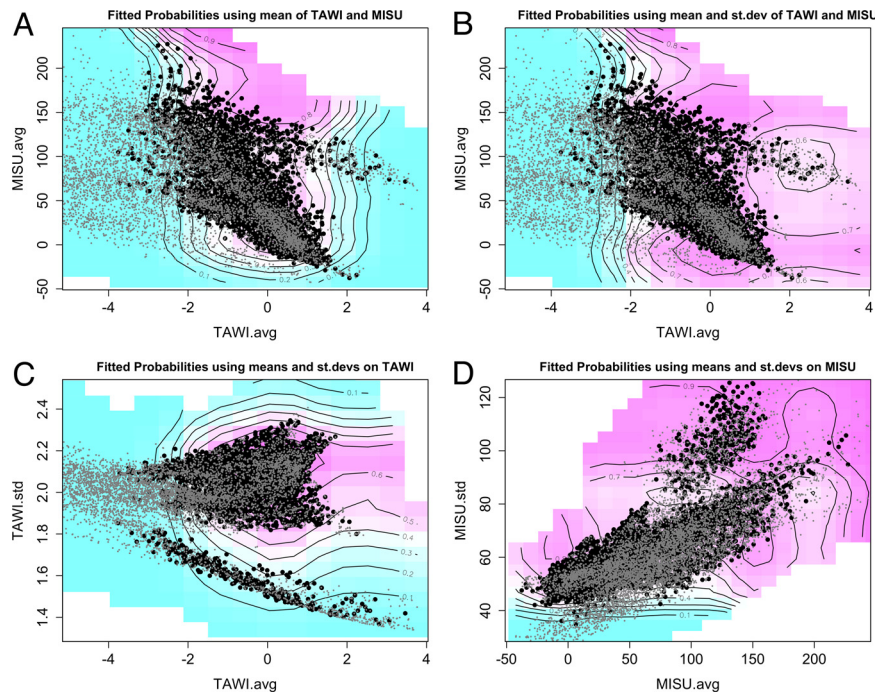


Fig. 4. Predicted probabilities for *F. sylvatica* along major climate gradients. (A and B) Illustration of fitted probabilities for *F. sylvatica* when using only mean climate predictors (A) compared with using means and standard deviations without statistical interaction term (B). (C and D) The probabilities from the model using means and standard deviations are plotted against TAWI.avg and TAWI.std (C) and against MISU.avg and MISU.std (D). Light gray and black dots represent all plots and plots where *F. sylvatica* is present, respectively. Blue and magenta represent low and high model probabilities, respectively, as indicated by the contour lines.

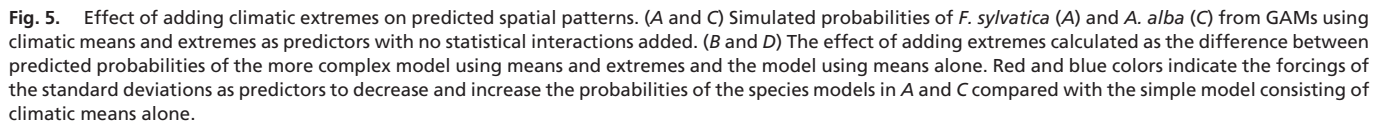
(Fig. S1 in *SI Appendix*). The overall improvement in cross-validated model performance was significant (TSS: interactions of variables representing means and extremes: 0.561 ± 0.034 ; mean only: -0.052 ± 0.038 ; $V = 0$, $P = 0.001$; AUC: interactions of mean and extreme variables: 0.848 ± 0.015 ; mean only: -0.030 ± 0.015 ; $V = 0$, $P = 0.001$). Adjusted D^2 increased by 20%, whereas TSS and AUC increased by 8% and 3%, respectively. However, when comparing the two versions of implementing extremes into models, we found that in 4 (*Acer pseudoplatanus*, *Pinus sylvestris*, *Quercus robur*, and *Betula pendula*) of the 11 species no significant improvement was reached in the ANOVA tests (Fig. S1 in *SI Appendix*). Only the overall improvement in cross-validated model accuracies was significant (TSS: interactions of mean and extreme variables: 0.561 ± 0.034 ; no interaction of mean and extreme variables: -0.012 ± 0.036 ; $V = 8$, $P = 0.024$; AUC: interactions of mean and extreme variables: 0.848 ± 0.015 ; no interaction of mean and extreme variables: -0.006 ± 0.016 ; $V = 9$, $P = 0.032$).

When adding variables representing extremes to models using climatic means for predicting spatial patterns of tree species, we note a shift in the probability space of *Fagus sylvatica* (Fig. 4). The species becomes less limited toward warmer temperatures and the limitation toward cold temperatures depends less on water availability (Fig. 4B). In the geographic space, adding variables representing extremes translates into a correction of predictions from the model using means only, especially in the dry and Mediterranean (Southern) parts of Switzerland (Fig. 5 A and B) for *F. sylvatica*. Including climate variability in the models of *Abies alba* (Fig. 5 C and D) increases prediction probabilities in the comparably moist climates of central and western Switzerland and further reduces the spatial distribution in the southern part of the Alps and in the central valleys with comparably dry climates. Presence/absence information for all tree species is given in Fig. S2 in *SI Appendix*; results for all 11 species are given in Fig. S3 in *SI Appendix*.

Discussion

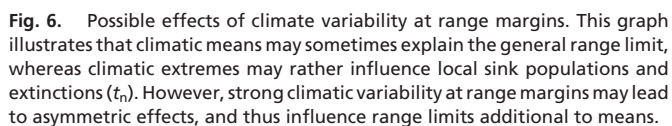
Our analyses reveal that complementing mean climate predictors with variables that represent climate extremes yields an improvement in the predictive power of species distribution models. The improvement is small compared with the model using climatic means only and specifically corrects spatial predictions compared with using climate means alone. This small improvement is partly in agreement with early investigations regarding explanations of distribution range patterns in trees and other plants. Larcher and Mair (47) observed that absolute climatic extremes alone are incapable of explaining the northern range limits of (climate constrained) Mediterranean oak species. They argued that average (winter) temperatures are more suitable for explaining northern range limits. We found, however, that adding climatic variability helps to explain such range limits. Another example involves the quest for a climatic explanation of the upper treeline worldwide. Current explanations focus on average climate predictors (48), and the best model based on a global dataset of mountain treeline temperatures reveals a strong relationship between treeline and average summer temperatures (49, 50). However, no explanation that we examined included additional effects of climate variability.

The effect of extremes and means are not easy to separate because of the high correlation between mean values and percentiles as measures of extremes. We expect that higher frequency or severity of extremes affect range margins in plants. Clear evidence exists that severe climate extremes influence plant demographic processes, such as growth (30, 32), regeneration (51), and mortality (38, 52, 53). However, it is not immediately evident from these studies that the observed responses affect range limits directly. Such processes may simply result in strong source-sink dynamics at range edges, with the range limit resulting from complex and multiple interactions (24, 54). For example, there may be strong mortality patterns after dry or cold years in tree saplings, but even a complete failure of



The association of range margin and climatic mean may not hold when climatic extremes occur with a skewed frequency distribution where the mean slowly shifts in a single direction, as during current, ongoing climatic change (57). In this situation mortality and regeneration are increasingly affected by climatic

The effects of climatic extremes on limiting species distributions are likely important when climate means shift geographically and when variability changes. The daily climate surfaces we used may have smoothed climatic extremes, causing us to overlook patterns of truly extreme events. The same could also occur because of the limited temporal span of the climate data. First, such limitations are difficult to overcome, because we used a contemporary method to generate daily climate surfaces. Second, we performed initial tests to use absolute extremes instead of means, which did not improve the model fits. Third, the size of the available time window to analyze extremes over a large area is potentially problematic. Trees are long-lived and their stress tolerance differs among life stages. More information on stress tolerance at different life stages would enhance our ability to optimize analyses to the relevant time window. Other effects that reduce the sensitivity of species distributions to climate fluctuations include human influence through forest management and the possibility that observed distributions may



not be in equilibrium with current climate (63). In our dataset, we assume a strong effect of forest management on the Swiss Plateau for *Picea abies*. For most forests of dry interior valleys we expect little to no effect from management on the elevational ranges of the species we considered. Finally, we believe that some of the limitations identified in our analysis of large-scale observational data can only be overcome by experiments. Ideally, such complementary analyses combine the power of large observational datasets with the insights of careful experimental design.

Materials and Methods

Tree Distribution Data. We used data from two forest inventory periods in Switzerland, which were sampled during the years 1983–1985 (NFI1) and 1995–1997 (NFI2) on a regular 1-km grid. In total, we had 12,328 plots available where species presence and absence for >50 species was derived from basal area estimates. The inventory records a total of >50 tree species, of which we selected 11 species for modeling. The selected species are (i) sufficiently abundant, (ii) belong to two different functional groups [broadleaf deciduous: *Fagus sylvatica* L., *Acer pseudoplatanus* L., *Fraxinus excelsior* L., *Quercus petraea* (Mattuschka) Lieblein, *Q. robur* L., *Sorbus aria* (L.) Crantz, *Betula pendula* Roth; needleleaf evergreen: *Picea abies* (L.) H. Karsten, *Abies alba* Miller, *Pinus sylvestris* L., *P. cembra* L.], and (iii) cover the full elevational gradient available in Switzerland from 180 m above seal level (a.s.l.) to the treeline situated at ≈2,450 m a.s.l. in the dry interior valleys. Additional details regarding the forest inventory data are given in *SI Appendix*.

Climate Predictor Data. We generated a climate predictor set containing long-term (1961–2006) averages of monthly, seasonal, or annual predictors and standard deviations of the mean values representing extremes. These estimates were derived at a spatial resolution of 100 m by using the DAYMET algorithm (64). We used data from automated weather stations (>180 for temperature; >350 for precipitation provided by MeteoSwiss) and a 100-m digital elevation model from the Swiss National Office of Topography. DAYMET generates output for daily average, minimum, and maximum temperature and precipitation. Additionally, we calculated potential evapotranspiration, and moisture index, which is the difference between precipitation and potential evapotranspiration (65). For the methods for deriving these two variables, see ref. 66. Additional to these climate predictors, we added slope and topographic position to the lists of evaluated predictors. More information on the derivation of the climate predictors is given in *SI Appendix*, and high and low mean and standard deviations of the selected climate variables are presented in Fig. S1 in *SI Appendix*.

Extreme climatic events are best analyzed by using specific approaches based on extreme value statistics (42). Frequencies of extreme events are, however, sensitive to the mean and even more to the variability of climatic variables at a given location (43, 44). Thus, adding extreme event indices as predictor variables to a model including the mean values would confound the effects of mean and variability. Therefore, we chose standard deviations as expressions of extremes in combination with means. We did this for three additional reasons, namely: (i) single-day or rare extremes are very difficult to assess or detect, and even more so to spatially extrapolate. Standard deviations that include the likelihood of extreme events are much easier to extrapolate and to measure. (ii) We do not know exactly what period is relevant

regarding extremes for each individual species. They are different in size and age, and the relevant period of sensitivity may change. Thus, a measure of variability (in combination with means) is more likely to capture the general likelihood of extreme events relevant to individual species than do single few observations. Variability can be calculated quite accurately from the whole time series. (iii) There is a high correlation between mean and extremes (percentiles). Thus, adding both means and percentiles as expressions of extremes would result in severe collinearity problems when fitting models.

For our model exercise with the selected tree species we chose two climatic variables, namely: (i) average winter temperature (TAWI: December, January, February), and (ii) average summer moisture index (MISU: June, July, August). TAWI (°C) expresses winter cold limitations, whereas MISU (mm) expresses water availability and levels of drought stress. For these two variables, we calculated both means and standard deviations. Additionally, we added slope and topographic position (66). The six selected variables show very low correlations on average (0.254), and only two variables correlate >0.5 (mean and SD of MISU: 0.73).

Statistical Analyses. We chose GAMs as implemented in the R package *mgcv* (67). This is a flexible, nonparametric method for calibrating the species response to topo-climatic predictors, which allowed us to additionally test the effect of interactions between means and standard deviations by using smooth terms built with tensor products (68). In *mgcv*, the degrees of smoothing are selected by internal cross-validations. All variables were entered in the default mode, and three models were finally calibrated for each species. The first used all selected variables except the two standard deviations, whereas the second included the two standard deviations. The third used interaction terms from tensor product smoothed terms of the respective mean and standard deviations (MISU and TAWI), instead of adding the two variables separately.

All models were 10-fold cross-validated and model performance of calibrated and cross-validated models was analyzed by calculating the adjusted D^2 (see ref. 5), threshold-maximized TSS (46) and AUC (69), which allows assessment of model accuracy independent of thresholds. The model improvement when adding extremes in addition to means was tested in two ways. First, we performed an ANOVA using a χ^2 test for checking the significance of the improvement in calibration strength. Second, we tested the improvement in model quality by applying a paired Wilcoxon test to the model evaluation values (TSS, AUC) of all species' models with means only against all species' models with means and extremes. The first (ANOVA) test allowed us to evaluate the individual improvement of models, whereas the second (Wilcoxon) test allowed us to evaluate the overall improvement of the predictive power of the models. All analyses were performed in the statistical environment R (70).

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