SUPPLEMENTARY INFORMATION FOR "Savannas of Asia: evidence for antiquity, current day biogeography and an uncertain future": Figures S1, S2, S3, Table S1, S2, S3 & Appendix 1

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Figure S1 Map of the Asia showing regions covered by this study. Regions shown are those defined by the Taxonomic Databases Working Group (TDWG) level 2 regions (Clayton et al 2016) with the exception of East Asia which is based on TDWG level 3 regions for southern China (Southeast China, Southwest China, Hainan Island). Key to colour codes for regions: Pink = South Asia (Indian subcontinent); Yellow = East Asia (southern China); Pale green = Continental Southeast Asia (Indochina); Dark green = Oceanic Southeast Asia (Malesia).

Figure S2 C₄ grass endemism and diversity in Asia as compared with other major savanna regions of the world. (a) C4 grass endemism per region versus total grass species per region. (b) Total C4 species per region as a function of region area in 1000s of km². Regions are all based on the Taxonomic Databases Working Group (TDWG) level 2 regions (Clayton et al. 2016) (see Fig. S1, Supplementary Materials) with the exception of East Asia which is based on level 3 regions (Southwest China, Southeast China, and Hainan Island). Key to codes for regions: $WIO = West Indian Ocean$; $SA = Southern Africa$; Aus = Australia; Br = Brazil; Ase1 = Indochina (continental Southeast Asia); Ase2 = Malesia (oceanic Southeast Asia); Ae = East Asia (southern China); As = India. See Table S1 for further details.

Figure S3 a) Observed distribution of savannas derived from vegetation maps for Africa (White 1983), Australia and South America (Lehmann et al 2011), and b) predicted distribution of savannas in the different continents using stochastic gradient boosting. The distribution of savannas was modeled separately for each continent based on climate, elevation and edaphic parameters, using the observed distribution map for that continent.

Table S1. Grass endemism and diversity as compared with other major savanna regions of the world calculated using TDWG regions (Clayton et al. 2016). Regions are all based on TDWG level 2 regions with the exception of East Asia, which is based on level 3 regions (see text). Size of regions were calculated using Eckert IV projections, following Vorontsova et al. 2016. Endemism of C4 grasses in the Asian TDWG regions (Fig S1, Supplementary Materials), was estimated using the GrassBase database (Clayton et al. 2016) combined with a database of C_3 / C4 pathways for grass taxa (Osborne et al. 2014).

Appendix 1: Methods for analysis of climate domains of Asian Savannas

Stochastic gradient boosting is an ensemble method for fitting statistical models that combines the strength of traditional statistical methods (decision trees) with machine learning techniques (boosting; Hastie et al 2001; Friedman et al. 2000, Friedman 2001, 2002). It works by iteratively building a number of small decision trees, each based on a random subset of the data, with each additional tree emphasizing observations poorly modeled by the existing collection of trees (Hastie et al 2001; Friedman et al. 2000, Lawrence et al. 2004, Elith et al. 2008). Finally, observations are assigned a class based on the most common classification amongst the trees (Lawrence et al. 2004, Elith et al. 2008). Gradient boosting is less sensitive to outliers and unbalanced data, is robust against overfitting, and has been shown to outperform many other classifiers (Friedman 2002, Lawrence et al. 2004).

We used existing vegetation maps to model the occurrence of savannas on different continents as a function of climate, elevation and edaphic parameters. For Africa, we used the continent-wide map developed by White (1983) to classify habitats as either savanna or non-savanna (see Sankaran et al. 2005), and for Australia and South America, we used the maps developed by Lehmann et al. (2011) (see Lehmann et al. 2011 for more details). Climate data were derived from the WorldClim climate database (Hijmans et al., 2005, http://www.worldclim.org), which provides data on 19 bioclimatic variables for the time period 1950-2000 (Hijmans et al., 2005), and soil nitrogen and percent clay from the ISRIC-WISE derived soil properties database (Baties 2012, www.isric.org). Potential evapotranspiration (PET) estimates were obtained from the Global Potential Evapo-Transpiration (Global-PET) database (http://www.cgiarcsi.org/data/global-aridity-and-pet-database; Zomer et al 2007, Zomer et al. 2008). All data were resampled to a resolution of 0.5° for our analysis.

For each continent, we first generated a training dataset by systematically sub-sampling every 4th pixel to account for issues of spatial autocorrelation. These training datasets were used to build continent-specific boosting models, which were then used to predict the distribution of savannas across the entire continent. Our training data set included 2547 pixels for Africa (52.6% of which were savanna), 750 for Australia (15.2% savanna) and 1625 for South America (9.9% savanna). We evaluated the accuracy of the different models based on the fraction of savanna pixels correctly identified. We evaluated two separate models, one including all 19 climate variables along with soil and elevation parameters as predictors, and the second which only included a subset of climatic predictor variables that were largely uncorrelated with one another along with soil and elevation parameters. Model performance did not differ consistently between the two, and we only report results here from the model built using the subset of climatic predictors. Our final set of predictor variables included mean annual temperature, annual temperature range, mean temperature of the driest quarter, mean annual precipitation, precipitation of the driest month, precipitation seasonality, potential evapotranspiration, soil N and clay contents, and elevation. Finally, we used the models developed for each continent to individually predict the potential distribution of savannas in Asia. Although our models were built using data sampled at a resolution of 0.5°, our predictions for Asia were based on climate and soil data sampled at resolution of 5 arc-minutes (approx. 9 km \times 9 km). All analyses were carried out using the 'caret' package (Liaw & Wiener, 2002) as implemented in R (R Core Team 2015).

Table S2. Model performance in terms of classification accuracies for Africa, Australia and South America.

Table S3. Area (in 1000 km²) of Asia predicted to support savannas as a function of prediction probabilities based on models developed for Africa, Australia and South America.

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