

Vegetation Community Composition and Species–Environment Relationships Along an Elevational Gradient in South-Central Bhutan

Abstract

Questions: Understanding how plant communities vary along elevation is essential for predicting biodiversity responses to environmental change. We asked how community composition, diversity, and species–environment relationships differ among vegetation strata along an elevational gradient, and which environmental variables best explain regeneration patterns.

Location: Sarpang District, south-central Bhutan; unmanaged subtropical to cool broadleaved forests spanning approximately 260 to 1 964 m a.s.l.

Methods: We analysed vegetation data from four strata (trees, shrubs, herbs, regeneration) across 220 plots. We constructed community matrices, quantified environmental variables, applied non-metric multidimensional scaling and canonical correspondence analysis, and tested group differences with permutational multivariate analysis of variance. Indicator species analysis and machine-learning models (random forest and gradient boosting) assessed regeneration patterns.

Results: Community composition differed among forest types in all strata (R^2 0.017 to 0.050; $p = 0.001$). Environmental predictors explained 3.2 to 3.8 percent of variation in trees and shrubs and were not significant for herbs and regeneration. Trees had the highest Shannon diversity (1.391 ± 0.595), whereas herbs had the lowest (0.325 ± 0.451). For regeneration richness, random forest showed modest cross-validated performance (root mean square error 1.165 ± 0.182 ; coefficient of determination 0.142 ± 0.040).

Conclusions: Forest types impose a statistically detectable but limited macro-environmental signal across strata. Community assembly is dominated by fine-scale heterogeneity and vertical decoupling among layers, and regeneration is only weakly predictable from coarse environmental gradients. These results provide a baseline for monitoring unmanaged broadleaved forests within the sampled 260–1 964 m range.

28 **Keywords:** elevational gradient, plant community, tropical mountain forest, species–
29 environment relationships, community assembly, diversity, regeneration, ordination, canonical
30 correspondence analysis, indicator species

31 **1 Introduction**

32 Elevational gradients constitute powerful natural laboratories for understanding how climate,
33 topography, and habitat heterogeneity interact to structure plant communities. Because tempera-
34 ture, moisture availability, and growing-season length vary predictably with altitude, mountain
35 systems have long served as test beds for ecological theory addressing species richness, envi-
36 ronmental filtering, and community turnover (Lomolino, 2001; Rahbek, 2005). Across regions,
37 richness–elevation relationships are frequently unimodal but vary with spatial extent, taxonomic
38 scope, and climatic context, indicating that no single mechanism universally governs elevational
39 diversity patterns (Grytnes and Vetaas, 2002; McCain, 2009). In forest ecosystems, steep abiotic
40 gradients combined with fine-scale variation in soils, canopy structure, and disturbance regimes
41 generate pronounced compositional turnover and vertically stratified responses among trees,
42 shrubs, and herb layers (Gilliam, 2007; Sundqvist et al., 2013). Mountain floras therefore often
43 exhibit high beta diversity over short distances, reflecting both dispersal limitation and niche
44 differentiation along climatic and edaphic gradients (Qian et al., 2005). Recent research has
45 increasingly focused on disentangling the relative roles of environmental filtering, functional
46 traits, and biotic interactions in shaping elevational community organization (Siefert et al., 2013;
47 Kraft et al., 2015). Nevertheless, many studies continue to focus primarily on canopy trees,
48 despite mounting evidence that understorey strata respond more sensitively to microclimatic and
49 edaphic variation and may follow distinct compositional trajectories along the same gradient
50 (Gilliam, 2007; Lenoir et al., 2008). Regeneration layers are especially informative in this
51 context because they integrate recent recruitment processes and may foreshadow future changes
52 in forest composition under ongoing climatic warming (Clark et al., 2014).

53 The eastern Himalaya represents a global centre of biodiversity characterized by extreme
54 elevational relief, complex terrain, and monsoonal climates that generate sharp forest transitions

55 and substantial species turnover (Myers et al., 2000; Grytnes and Vetaas, 2002). Within this
56 region, numerous studies have documented steep elevational zonation and marked diversity
57 patterns, while broad-scale syntheses emphasize the region's sharp biodiversity gradients and
58 habitat heterogeneity (Myers et al., 2000; Grytnes and Vetaas, 2002). Despite this growing body
59 of work, community-level investigations integrating multiple vegetation strata across Himalayan
60 elevational gradients remain comparatively uncommon. Few studies explicitly compare species–
61 environment relationships among canopy and understorey layers, and regeneration dynamics
62 are seldom analysed in parallel with diversity and compositional patterns, despite their central
63 importance for forest resilience and long-term reorganization (Sundqvist et al., 2013; Clark et al.,
64 2014).

65 Here, we present a multi-stratum assessment of plant community composition, diversity,
66 and species–environment relationships along an elevational gradient in south-central Bhutan.
67 We test whether forest types differ in composition across four strata—trees, shrubs, herbs, and
68 regeneration—identify macro-environmental correlates of community structure, and quantify
69 elevational trends in alpha and beta diversity together with indicator species. We also use
70 machine-learning models as exploratory tools to evaluate which measured variables are asso-
71 ciated with regeneration richness. By integrating multiple vegetation strata within a unified
72 framework, this study provides one of few multi-stratum assessments in the eastern Himalaya
73 and explicitly evaluates scale mismatch between macro predictors and microsite processes.

74 **2 Methods**

75 **2.1 Study area**

76 The study was conducted in unmanaged forests of Sarpang District in south-central Bhutan under
77 the jurisdiction of Divisional Forest Office, Sarpang. The region spans a pronounced elevational
78 gradient from approximately 153 to 3 500 m a.s.l., encompassing subtropical broadleaved forests
79 at lower elevations (100–500 m), warm broadleaved forests at mid-elevations (500–1 500 m),
80 and cool broadleaved forests between 1 500 and 3 000 m. Such steep altitudinal transitions typify
81 Himalayan landscapes and generate strong climatic and edaphic gradients over short distances

82 (Grytnes and Vetaas, 2002; Körner, 2007) (Figure 1).

83 This broader regional range provides ecological context; all statistical inference in this
84 study is based on sampled plots spanning approximately 260 to 1 964 m a.s.l.

85 Annual precipitation is high and strongly seasonal, reflecting monsoon circulation inten-
86 sified by orographic uplift, a defining feature of the eastern Himalayan foothills (Bookhagen and
87 Burbank, 2010). Rugged terrain and variable slope exposure create pronounced microclimatic
88 heterogeneity, which is expected to contribute to high beta diversity and fine-scale community
89 turnover along the gradient (Grytnes and Vetaas, 2002). The landscape also forms part of
90 a strategic ecological linkage among Bhutan's protected areas, although the sampled forests
91 themselves lie largely outside formal management regimes.

92 **2.2 Sampling design**

93 Vegetation sampling followed a stratified random design along the elevational gradient extending
94 from Shershong (approximately 260 m a.s.l.) in the south to Singye (approximately 1 964 m a.s.l.)
95 in the north. A total of 220 plots were established across forest types to ensure representation of
96 major vegetation zones and environmental conditions.

97 Each plot measured 20×20 m (400 m^2), a size widely used in forest community studies
98 to characterise tree assemblages and stand structure, with a nested 2×2 m subplot used to
99 survey herbaceous vegetation. Fieldwork was undertaken between March and November to
100 encompass seasonal variability in understorey communities while avoiding periods of peak
101 monsoon inaccessibility. Plots were distributed approximately evenly among elevational bands
102 and forest types to ensure balanced representation of vegetation zones. Within each band, plot
103 locations were stratified by forest type and positioned using random bearings and distances from
104 access transects, subject to terrain safety and forest accessibility constraints.

105 **2.3 Vegetation data collection**

106 All free-standing woody individuals with diameters at breast height (DBH) > 10 cm and height
107 > 1.3 m were recorded as trees. DBH was measured using diameter tapes, and total height was
108 estimated by using hypsometers. Shrubs were assessed for presence, height, and lateral spread,

109 while herbaceous species were quantified within subplots using percentage cover, frequency,
110 and maximum height.

111 Regeneration comprised all woody individuals below the canopy layer that had not yet
112 reached the tree stratum. Seedlings were defined as individuals ≤ 1.3 m in height, irrespective
113 of stem diameter, whereas saplings were defined as individuals > 1.3 m in height with DBH \leq
114 10 cm. Species exceeding 10 cm DBH were assigned to the tree layer. Regeneration richness
115 therefore integrated both seedling and sapling cohorts, representing early demographic stages
116 that are particularly sensitive to microsite conditions and neighborhood effects.

117 Species identifications were verified using regional floras and herbarium reference mate-
118 rial, and taxonomic nomenclature was cross-checked against the World Flora Online database
119 (accessed 6 Feb 2026). Structural attributes, including basal area, DBH-class distributions,
120 and height-class distributions, were derived to characterize stand structure and regeneration
121 dynamics across forest types and elevational gradients.

122 **2.4 Environmental variables**

123 Topographic attributes (latitude and longitude), including elevation, slope, and aspect, were
124 recorded in the field using handheld Global Positioning System receivers, clinometers, and com-
125 passes. Spatial climatic surfaces describing air temperature, precipitation, evapotranspiration,
126 and water balance were extracted from the national gridded climate dataset produced by Dorji
127 et al. (2025), developed by CSIRO and distributed with a permanent digital object identifier
128 (<https://doi.org/10.25919/pec2-hs50>).

129 Environmental values were extracted at plot coordinates using the dataset's native grid
130 and historical baseline layers (1986–2015), consistent with the metadata provided for the
131 archived climate product.

132 These climate layers were used consistently across all spatial and statistical analyses.

133 **2.5 Data preparation**

134 Field records were curated in tabular format to compute species abundance, frequency, basal
135 area and importance value index separately for each vegetation stratum. Alpha diversity was

136 quantified using the Shannon–Wiener index (H'), Simpson’s diversity index ($1 - D$) and Pielou’s
137 evenness (J) for each forest type and vegetation layer (Shannon, 1948; Simpson, 1949; Pielou,
138 1966).

139 **2.6 Multivariate community analyses**

140 Floristic variation among plots was examined using non-metric multidimensional scaling
141 (NMDS) based on Bray–Curtis dissimilarities (Bray and Curtis, 1957; Kruskal, 1964). Species–
142 environment relationships were assessed using canonical correspondence analysis (CCA), which
143 constrains ordination axes by linear combinations of climatic and topographic predictors (ter
144 Braak, 1986). Ordination axes were interpreted through correlations between site scores and
145 environmental variables.

146 Community differences among forest types were tested using permutational multivariate
147 analysis of variance (Anderson, 2001). NMDS was computed with two dimensions ($k = 2$)
148 and a fixed random seed (42) to ensure reproducibility. Environmental vectors were fitted
149 using permutation tests with 999 randomizations. PERMANOVA was conducted with 999
150 permutations, and homogeneity of multivariate dispersions was evaluated using permutation
151 tests on distance-to-centroid values (Warton et al., 2012).

152 To support valid interpretation, PERMANOVA results were interpreted jointly with
153 effect sizes (R^2) and dispersion tests. In strata with heterogeneous dispersion, PERMANOVA
154 was interpreted as reflecting combined centroid and spread effects rather than pure location shifts
155 (Warton et al., 2012). NMDS was used for pattern visualization rather than as a stand-alone test
156 of group separation.

157 Prior to CCA, collinearity among predictors was assessed using variance inflation factors,
158 and variables exceeding a threshold of 10 were sequentially removed following deterministic se-
159 lection rules. Statistical significance of constrained axes and individual predictors was evaluated
160 using 999 permutations.

161 Indicator species analysis was applied to identify taxa significantly associated with
162 forest types using the indicator value method, with significance assessed at $\alpha = 0.05$ and 999
163 permutations (Dufrêne and Legendre, 1997).

164 **2.7 Regeneration dynamics and predictive modelling**

165 Seedling and sapling data were analysed to quantify regeneration patterns and spatial structure
166 across forest types and elevations. Environmental drivers of regeneration richness were modelled
167 using random forest and extreme gradient boosting algorithms, both of which are widely applied
168 for non-linear ecological prediction problems (Breiman, 2001; Chen and Guestrin, 2016).
169 Random forest models were fitted with 500 trees, and predictor importance was quantified as
170 the percentage increase in mean squared error following permutation. Gradient boosting models
171 were parameterized with 100 boosting rounds, a maximum tree depth of four and a learning rate
172 of 0.1, and predictor influence was assessed using relative gain values.

173 Model performance was evaluated using five-fold cross-validation with a fixed random
174 seed (42) to ensure reproducibility. Predictive accuracy was summarized using cross-validated
175 error statistics and explained variance, and response curves were inspected to verify ecologically
176 plausible relationships between regeneration patterns and key climatic or topographic predictors.

177 Because predictive performance was modest, variable-importance results were inter-
178 preted as exploratory indicators of potential drivers rather than definitive causal rankings.

179 **2.8 Software environment**

180 All analyses were conducted in R. Community-ecology workflows, including ordination,
181 dissimilarity-based testing and indicator-species analysis, were implemented primarily using
182 functions described in the ecological literature underlying widely adopted multivariate methods
183 (ter Braak, 1986; Anderson, 2001; Dufrêne and Legendre, 1997). Results from selected multi-
184 variate classifications and indicator-species procedures were independently verified in PC-ORD
185 version 5 to confirm analytical consistency. Microsoft Excel was used for preliminary data
186 curation and quality-control screening prior to statistical analysis.

187 **2.9 Reproducibility and data stewardship**

188 Reproducibility was ensured through systematic archiving of raw field observations, spatial
189 environmental layers, curated analysis tables and derived outputs. All distance measures, ordina-

190 tion settings, permutation schemes and machine-learning hyperparameters were documented in
191 metadata files accompanying the archived datasets. Intermediate products, including ordination
192 scores, regeneration prediction surfaces, cross-validation partitions and variable-importance
193 tables were retained to permit independent replication of all analytical steps.

194 Upon submission, data and code supporting the results will be deposited in a public repos-
195 itory with a digital object identifier in accordance with data-sharing policy, with anonymized
196 access provided during double-blind peer review.

197 **3 Results**

198 **3.1 Community composition and ordination**

199 Non-metric multidimensional scaling (NMDS) showed modest but detectable compositional
200 differentiation among forest types across strata (Figure 2). Stress values were uniformly low
201 (trees = 0.0001; shrubs = 0.070; herbs = 0.001; regeneration = 0.0006), indicating a low-stress
202 two-dimensional rank-order fit of Bray–Curtis dissimilarities rather than a perfect ordination (Ap-
203 pendix S3). We explored multiple dimensional solutions, and stress values remained consistently
204 low, confirming ordination stability.

205 The near-zero stress values for trees, herbs, and regeneration likely arise from matrix
206 properties that ease low-dimensional rank fitting (for example, high tied dissimilarities and sparse
207 compositional structure), so ordinations were interpreted conservatively as broad summaries
208 rather than precise geometric distances (Kruskal, 1964; Clarke, 1993). We therefore cross-
209 checked NMDS patterns against PERMANOVA, dispersion tests, and constrained ordination
210 instead of relying on visual separation alone.

211 Shrub and herb assemblages displayed the greatest dispersion in ordination space and the
212 strongest alignment with fitted climatic vectors, particularly temperature, evapotranspiration and
213 precipitation (Figure 2B–C). Tree assemblages were comparatively compressed along NMDS
214 axes (Figure 2A), while regeneration plots showed broad scatter (Figure 2D), together indicating
215 vertical decoupling and strong neighbourhood-scale heterogeneity.

216 These stratified responses mirror documented altitudinal forest-type transitions along

217 Bhutanese dry valley slopes (Wangda and Ohsawa, 2006a).

218 **3.2 Community differences among forest types**

219 Community composition differed significantly among forest types across all vegetation strata (Ta-
220 ble 1). Tree assemblages exhibited modest but statistically significant separation (PERMANOVA
221 $R^2 = 0.031$, $F = 3.38$, $p = 0.001$), with homogeneous dispersions (betadisper $p = 0.291$), indi-
222 cating that group differences reflected centroid shifts rather than unequal within-group variability
223 (Appendix S2).

224 Shrub communities showed the strongest differentiation among forest types ($R^2 = 0.050$,
225 $F = 5.05$, $p = 0.001$), although dispersions were heterogeneous ($p = 0.001$), suggesting that
226 both location and spread contributed to observed differences. Herb and regeneration assemblages
227 also differed significantly ($R^2 = 0.017$ and 0.021 , respectively; both $p = 0.001$), but only the
228 herb stratum showed heterogeneous dispersion (Table 1).

229 The greater sensitivity of shrub and herb layers relative to trees is consistent with reported
230 altitudinal zonation and vegetation transitions in Himalayan forests (Wangda and Ohsawa, 2006a;
231 Grierson and Long, 2001). In strata with heterogeneous dispersion, PERMANOVA reflects
232 combined centroid and spread effects. Accordingly, shrub and herb results were interpreted as
233 mixed location–dispersion signals, whereas homogeneous dispersions for trees and regeneration
234 indicate that their compositional differences more closely represent centroid shifts.

235 **3.3 Species–environment relationships**

236 Canonical correspondence analysis revealed that selected environmental variables—aspect,
237 evapotranspiration, latitude, longitude and slope—explained a small but statistically significant
238 fraction of constrained inertia in trees and shrubs. The percentage of total inertia explained
239 was 3.2% for trees ($p = 0.008$) and 3.8% for shrubs ($p = 0.001$; Table 2; Figure 5). For herbs
240 and regeneration, constrained models explained 3.4% of total inertia but were not significant at
241 $\alpha = 0.05$ ($p = 0.125$ and 0.076 , respectively) (Appendix S9).

242 CCA biplots illustrated contrasting environmental associations among strata (Figure 5).
243 Tree communities aligned primarily with slope and aspect gradients, shrub assemblages were

244 most strongly oriented toward evapotranspiration and latitudinal position, while herbaceous
245 vegetation responded to combined slope–climate axes. Regeneration patterns were weakly
246 associated with evapotranspiration and aspect.

247 The low proportion of constrained inertia across all strata is expected in steep, hetero-
248 geneous Himalayan terrain and is consistent with substantial unmeasured microsite drivers
249 operating within broader altitudinal transitions (Grytnes and Vetaas, 2002; Wangda and Ohsawa,
250 2006a).

251 **3.4 Diversity patterns**

252 Alpha diversity differed markedly among vegetation strata (Table 3; Figure 3). Trees exhibited
253 the highest mean species richness (5.30 ± 2.57 species per plot) and Shannon diversity ($1.391 \pm$
254 0.595), followed by shrubs (richness = 4.77 ± 3.43 ; Shannon = 1.063 ± 0.706), regeneration
255 (richness = 2.00 ± 1.26 ; Shannon = 0.443 ± 0.499) and herbs (richness = 1.79 ± 1.32 ; Shannon
256 = 0.325 ± 0.451). Pielou’s evenness was highest for trees (0.903) and lowest for herbs (0.804).

257 Species accumulation curves approached asymptotes most rapidly for shrubs and regen-
258 eration, whereas trees and herbs continued to increase steadily with sampling effort, indicating
259 incomplete saturation of the regional species pool in these strata (Figure 4). Such patterns are
260 consistent with documented altitudinal zonation of Bhutanese dry-valley forests and reported
261 regeneration dynamics of dominant trees (Wangda and Ohsawa, 2006a,b). Continued species
262 accumulation in tree and herb strata suggests that additional sampling would likely yield further
263 taxa, particularly rare canopy species and ephemeral understorey herbs, indicating substantial
264 compositional turnover.

265 Species richness varied along the elevational gradient for all strata (Figure 7). Shrubs
266 displayed a unimodal mid-elevation peak, whereas trees showed comparatively stable richness
267 across the sampled range, and herbs and regeneration exhibited weaker, irregular trends. These
268 responses align with Himalayan studies reporting mid-elevation richness peaks and altitudinal
269 zonation in woody layers (Acharya et al., 2011; Grytnes and Vetaas, 2002; Wangda and Ohsawa,
270 2006a).

271 Beta diversity (Whittaker’s β) was highest for herbs (75.04), followed by regeneration

272 (54.63), trees (41.86) and shrubs (23.89), indicating pronounced species turnover in the herb
273 layer relative to the regional species pool.

274 **3.5 Indicator species**

275 Indicator-species analysis detected significant associations with elevational classes across all
276 strata (Table 4). Shrubs contained the largest proportion of indicator taxa (15 of 101 species;
277 14.9%), followed by herbs (10 of 134; 7.5%), regeneration (8 of 109; 7.3%) and trees (10 of
278 221; 4.5%) (Appendix S4).

279 Among canopy trees, *Schima wallichii* and *Castanopsis indica* were characteristic of
280 low- to mid-elevation forests, whereas *Alnus nepalensis* was associated with higher elevations.
281 These patterns accord with floristic zonation documented in Bhutanese mountain forests, where
282 evergreen broadleaved species dominate lower belts and pioneer or cool-adapted taxa become
283 more frequent upslope (Wangda and Ohsawa, 2006a; Grierson and Long, 2001).

284 **3.6 Regeneration modelling**

285 Random forest produced modest predictive performance for regeneration richness (Table 5;
286 Figure 6), with cross-validated $R^2 = 0.142 \pm 0.040$ and RMSE 1.165 ± 0.182 . Gradient boosting
287 showed near-zero to negative explanatory power ($R^2 = -0.009 \pm 0.096$; RMSE = 1.261 ± 0.192)
288 (Appendix S5).

289 Fold-level diagnostics (Appendix S7) showed moderate variability in RF performance
290 (R^2 range: 0.096–0.193; RMSE range: 0.999–1.476) and unstable XGBoost performance (R^2
291 range: –0.154 to 0.106; RMSE range: 1.102–1.570), reinforcing that coarse predictors provide
292 only weak predictability of recruitment.

293 Variable-importance rankings indicated that shrub-layer attributes, together with temper-
294 ature, elevation and evapotranspiration, were the strongest correlates of regeneration richness
295 (Figure 6; Appendix S6). Ecologically, this pattern is consistent with neighbourhood-scale
296 filtering and local biotic interactions outweighing coarse macro-environmental gradients for
297 early life stages (Wangda and Ohsawa, 2006b).

298 Given the low explanatory power, these models are interpreted as exploratory ecological

299 inference rather than deterministic prediction. Important drivers—such as canopy openness,
300 disturbance history, soil properties or fine-scale moisture regimes—were not fully captured
301 by the available predictors, a common limitation in forest-regeneration modelling in complex
302 mountain terrain (Elith et al., 2008).

303 **4 Discussion**

304 **4.1 Stratified community turnover along elevational gradients**

305 Our analyses indicate clear vertical stratification in community turnover, with understorey as-
306 semblages (shrubs and herbs) more variable than canopy trees. This pattern mirrors observations
307 from other Himalayan forests where understorey communities respond strongly to microclimatic
308 and edaphic heterogeneity, whereas long-lived canopy trees integrate conditions over broader
309 temporal and spatial scales (Wangda and Ohsawa, 2006a; Grytnes and Vetaas, 2002). Forest-type
310 effects were statistically detectable but limited in magnitude: PERMANOVA explained only
311 1.7–5.0% of compositional variation ($R^2 = 0.017\text{--}0.050$, $p = 0.001$). These low effect sizes
312 are compatible with high beta diversity and fine-grained heterogeneity in montane systems
313 (Acharya et al., 2011; Carpenter, 2005). Importantly, interpretation depends on dispersion
314 structure: in shrub and herb strata, heterogeneous dispersions indicate mixed location–spread
315 effects, whereas tree and regeneration strata with homogeneous dispersions indicate clearer
316 centroid shifts among forest types. This cross-stratum contrast supports vertical decoupling,
317 where canopy differentiation is detectable but modest and understorey turnover is dominated by
318 neighbourhood-scale heterogeneity.

319 **4.2 Environmental drivers and scale of control**

320 Direct gradient analyses (CCA) showed that aspect, slope, evapotranspiration, and spatial
321 position explained only a small fraction of floristic variation ($\approx 3\text{--}4\%$ in trees and shrubs).
322 Rather than indicating weak study design, this low constrained inertia is expected in steep
323 Himalayan terrain, where microtopography, canopy gaps, soils, and disturbance history vary
324 at much finer scales than gridded macro predictors (Grytnes and Vetaas, 2002; Wangda and

325 Ohsawa, 2006a). The contrasting orientations among strata reinforce this scale mismatch.
326 Tree communities aligned mainly with slope and aspect, while shrubs and regeneration were
327 more associated with evapotranspiration and latitude; herbs tracked mixed slope–climate axes.
328 Together, these patterns indicate that macro-environmental gradients are detectable but limited,
329 and that community assembly is dominated by microsite processes and vertical decoupling
330 among strata.

331 **4.3 Diversity patterns and beta-diversity**

332 Marked differences in alpha diversity among strata further highlight the vertically structured
333 nature of Himalayan broadleaved forests. Tree layers in our study maintained high species
334 richness and evenness across the gradient, reflecting the coexistence of multiple canopy domi-
335 nants in these unmanaged stands – a feature repeatedly documented in Bhutan’s warm and cool
336 broadleaved forests (Grierson and Long, 2001; Wangda and Ohsawa, 2006a). Shrub richness
337 peaked at mid-elevations, consistent with many Himalayan studies that find unimodal diversity
338 curves in the woody understorey around intermediate altitudes where moisture, temperature,
339 and structural complexity are jointly optimized (Grytnes and Vetaas, 2002). By contrast, herbs
340 and regeneration showed weak or irregular elevational trends in richness, coupled with elevated
341 beta diversity relative to alpha diversity in the herb layer. This pattern implies that small-scale
342 environmental mosaics, rather than smooth elevational forcing, structure ground-layer com-
343 munities. High herb-layer beta diversity is widely reported in Himalayan forests and alpine
344 systems, where aspect, litter depth, and gap dynamics generate sharp species turnover over short
345 distances (Grytnes and Vetaas, 2002). The fact that our species accumulation curves did not
346 fully asymptote for trees and herbs even after 221 plots suggests that a considerable portion
347 of the regional species pool remained unsampled in these strata, consistent with substantial
348 compositional turnover.

349 **4.4 Indicator species and floristic transitions**

350 Indicator species analysis revealed fewer significant canopy indicators than shrub and herb
351 indicators, supporting vertical decoupling in habitat filtering strength. Low tree indicator

352 specificity is consistent with broader niche amplitudes in canopy taxa across the sampled gradient.
353 In contrast, stronger shrub and herb differentiation implies finer-scale filtering in understorey
354 microsites. This contrast aligns with Himalayan zonation as overlapping distributions rather
355 than discrete boundaries (Ohsawa, 1995; Wangda and Ohsawa, 2006a).

356 **4.5 Regeneration dynamics and model interpretation**

357 Machine-learning results were used as exploratory ecological evidence on regeneration con-
358 trols. The best model (random forest) explained only about 14% of cross-validated variance
359 ($R^2 \approx 0.14$), indicating weak predictability from the available coarse predictors. Shrub-layer
360 structure emerged as the strongest correlate, supporting neighbourhood-scale filtering during
361 early recruitment. This interpretation is consistent with Bhutanese forest observations that local
362 stand structure influences regeneration (Wangda and Ohsawa, 2006a,b). The low predictive
363 power also indicates missing microsite drivers, such as canopy openness, soil properties, distur-
364 bance history, and fine-scale moisture (Elith et al., 2008). Accordingly, these models should not
365 be read as deterministic forecasts, but as evidence that regeneration is only weakly predictable
366 from macro-environmental gradients in this landscape.

367 **4.6 Limitations and novelty of the study**

368 Several caveats merit consideration. First, climate variables were derived from interpolated
369 surfaces (Dorji et al., 2025) and may not capture plot-level microclimate in steep terrain. Second,
370 heterogeneous dispersion in shrub and herb strata means that part of PERMANOVA significance
371 reflects spread differences, not only centroid shifts. Third, regeneration measurements represent
372 a temporal snapshot and may miss episodic establishment dynamics of slow-growing montane
373 trees. Spatial autocorrelation was not explicitly modeled, which may influence fine-scale infer-
374 ence, and future work could apply spatially explicit approaches. Even with these constraints,
375 this study provides one of few multi-stratum assessments in the eastern Himalaya. The main
376 contribution is to show that broad-scale environmental effects are statistically detectable but lim-
377 ited, while fine-scale heterogeneity, neighbourhood filtering, and vertical decoupling dominate
378 assembly patterns.

5 Conclusions

Across 260–1 964 m a.s.l. in south-central Bhutan, forest types imposed a statistically detectable but limited macro-environmental signal on community composition ($R^2 = 0.017\text{--}0.050$). Understorey strata showed greater turnover and higher dispersion than canopy trees, indicating strong vertical stratification and decoupling among layers.

CCA explained only about 3–4% of variation, which is consistent with steep Himalayan terrain where microsite heterogeneity is high and many local drivers are unmeasured. PERMANOVA interpretation was therefore conditioned on dispersion structure, with mixed centroid–spread effects in heterogeneous strata.

Regeneration was only weakly predictable from coarse predictors (random forest CV $R^2 \approx 0.14$), and shrub-layer attributes emerged as the strongest correlates. This supports neighbourhood-scale filtering during early recruitment and reinforces that macro gradients alone do not capture regeneration dynamics.

This study provides a baseline for long-term monitoring of unmanaged broadleaved forests in the sampled elevation range. Inference should remain restricted to this 260–1 964 m gradient.

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Table 1: Results of permutational multivariate analysis of variance (PERMANOVA) and tests of multivariate dispersion for vegetation strata across forest types. p -values are based on 999 permutations.

| Stratum | Sites | Groups | R^2 | F | p | Disp. F | Disp. p | Homogeneous |
|--------------|-------|--------|-------|------|-------|-----------|-----------|-------------|
| Trees | 216 | 3 | 0.031 | 3.38 | 0.001 | 1.24 | 0.291 | Yes |
| Shrubs | 194 | 3 | 0.050 | 5.05 | 0.001 | 12.93 | 0.001 | No |
| Herbs | 206 | 3 | 0.017 | 1.70 | 0.001 | 15.05 | 0.001 | No |
| Regeneration | 205 | 3 | 0.021 | 2.14 | 0.001 | 1.17 | 0.319 | Yes |

Table 2: Summary of canonical correspondence analyses by vegetation stratum. Environmental variables were selected following iterative variance inflation factor screening (threshold = 10). Significance of constrained inertia was assessed using 999 permutations.

| Stratum | Sites | Species | Env. vars | Total inertia | Constrained | % Explained | p |
|--------------|-------|---------|-----------|---------------|-------------|-------------|-------|
| Trees | 216 | 221 | 5 | 35.117 | 1.109 | 3.2 | 0.008 |
| Shrubs | 194 | 101 | 5 | 27.832 | 1.045 | 3.8 | 0.001 |
| Herbs | 206 | 134 | 5 | 68.475 | 2.309 | 3.4 | 0.125 |
| Regeneration | 205 | 109 | 5 | 58.906 | 1.987 | 3.4 | 0.076 |

Table 3: Mean alpha-diversity metrics by vegetation stratum. Values are mean \pm standard deviation.

| Stratum | Plots | Richness | Shannon | Simpson | Evenness |
|--------------|-------|-----------------|-------------------|---------|----------|
| Trees | 221 | 5.30 \pm 2.57 | 1.391 \pm 0.595 | 0.665 | 0.903 |
| Shrubs | 198 | 4.77 \pm 3.43 | 1.063 \pm 0.706 | 0.524 | 0.832 |
| Regeneration | 209 | 2.00 \pm 1.26 | 0.443 \pm 0.499 | 0.264 | 0.839 |
| Herbs | 210 | 1.79 \pm 1.32 | 0.325 \pm 0.451 | 0.193 | 0.804 |

Table 4: Summary of indicator-species analysis by stratum. Groups were defined by elevational class (low, mid and high). Significance was assessed using 999 permutations at $\alpha = 0.05$.

| Stratum | Sites | Species | Groups | Significant indicators | % Significant |
|--------------|-------|---------|--------|------------------------|---------------|
| Trees | 216 | 221 | 3 | 10 | 4.5 |
| Shrubs | 194 | 101 | 3 | 15 | 14.9 |
| Herbs | 206 | 134 | 3 | 10 | 7.5 |
| Regeneration | 205 | 109 | 3 | 8 | 7.3 |

Table 5: Cross-validated performance of machine-learning models predicting regeneration richness across five folds. R^2 denotes coefficient of determination; RMSE, root mean square error; MAE, mean absolute error. Full fold-level output is provided in Appendix S7.

| Model | Predictors | Obs. | CV R^2 | CV RMSE | CV MAE |
|---------------|------------|------|--------------------|-------------------|--------|
| Random Forest | 13 | 192 | 0.142 ± 0.040 | 1.165 ± 0.182 | 0.907 |
| XGBoost | 13 | 192 | -0.009 ± 0.096 | 1.261 ± 0.192 | 0.942 |

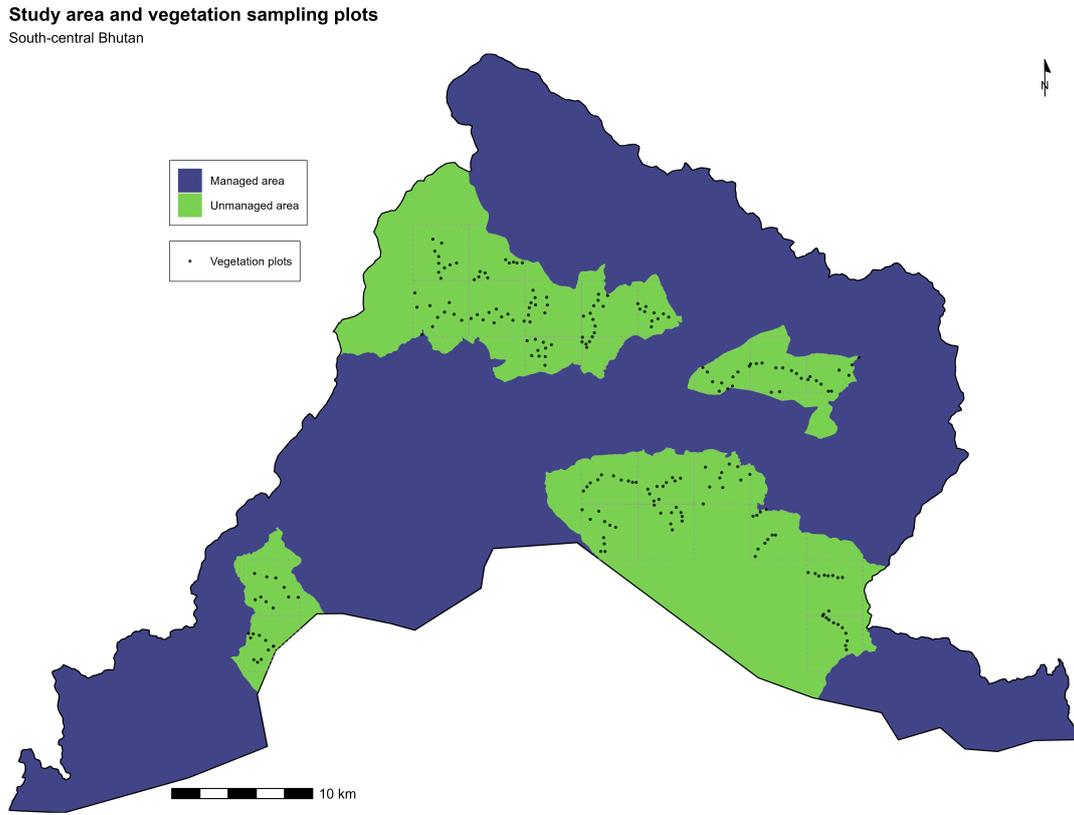


Figure 1: Study area and vegetation sampling plots in south-central Bhutan (Sarpang District). Green polygons indicate unmanaged forest areas and black points represent sampled vegetation plots along the elevational gradient.

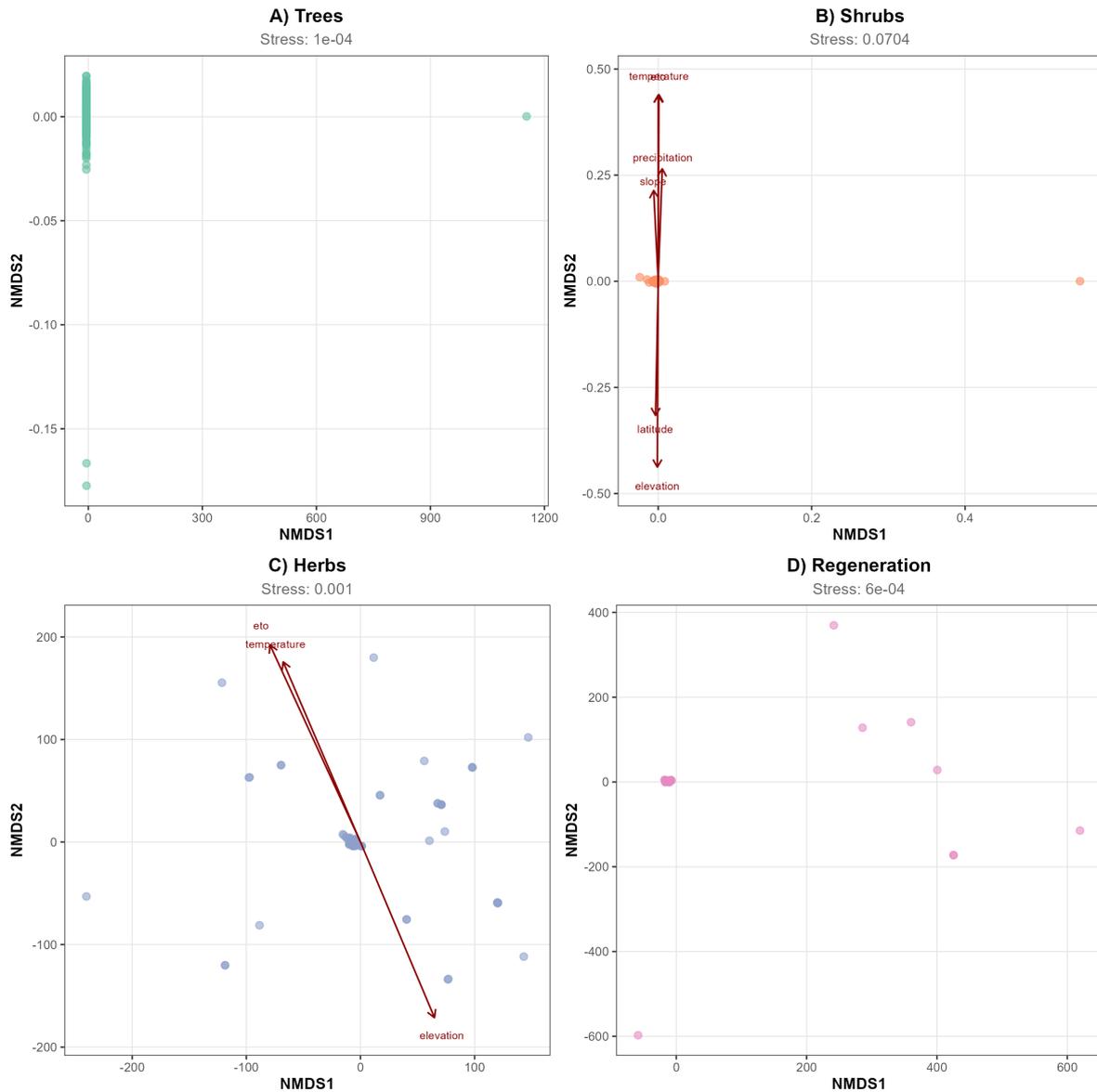


Figure 2: Non-metric multidimensional scaling (NMDS) ordinations of vegetation plots for four strata: (A) trees, (B) shrubs, (C) herbs and (D) regeneration. Points represent sample plots coloured by forest type; polygons delineate convex hulls for each forest type. Red arrows indicate fitted environmental vectors significant at $p < 0.05$ based on 999 permutations. Stress values are shown for each ordination.

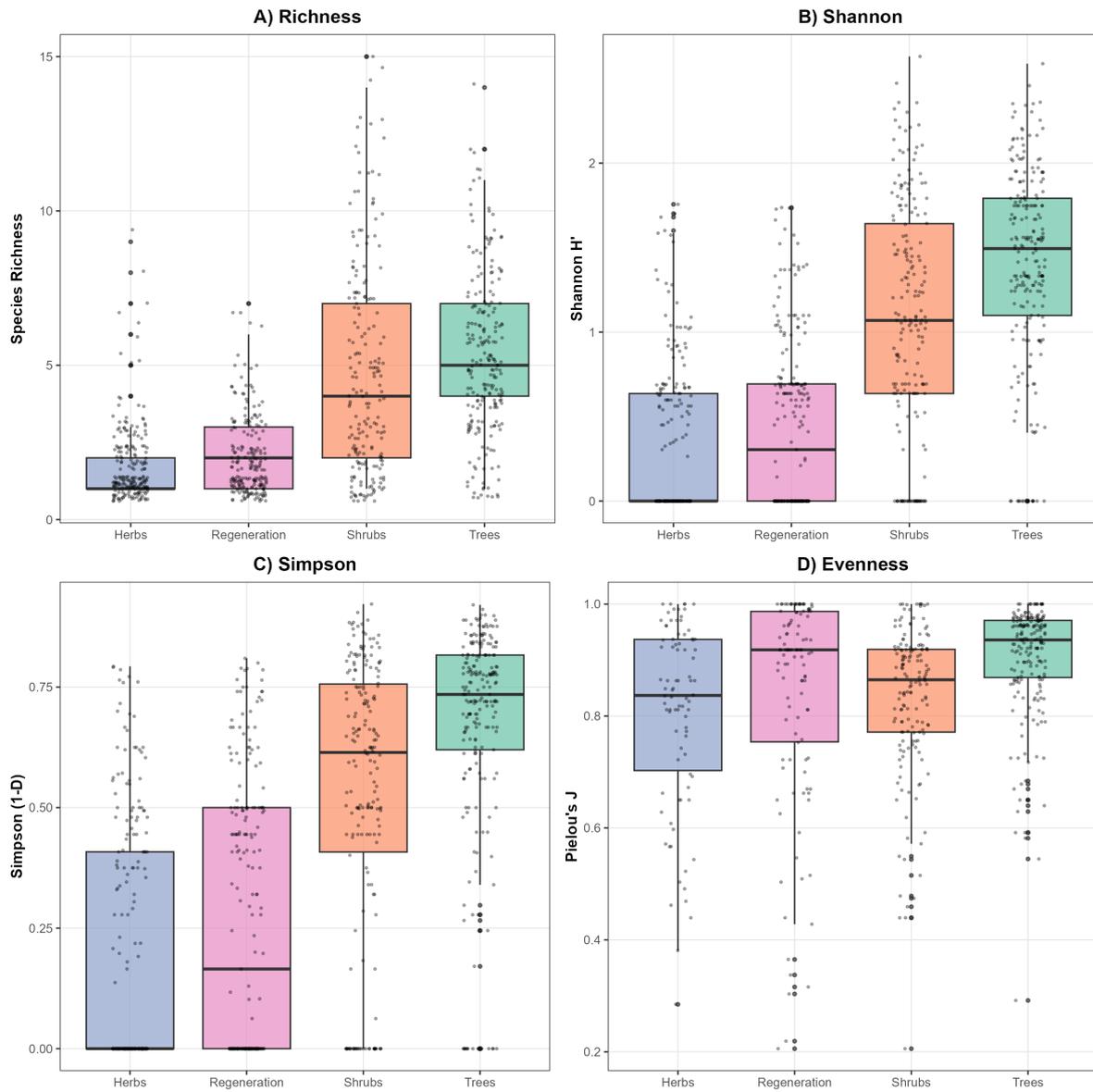


Figure 3: Alpha-diversity metrics across vegetation strata. Panels show (A) species richness, (B) Shannon diversity (H'), (C) Simpson diversity ($1 - D$) and (D) Pielou's evenness (J). Boxplots depict medians and interquartile ranges; points represent individual plots.

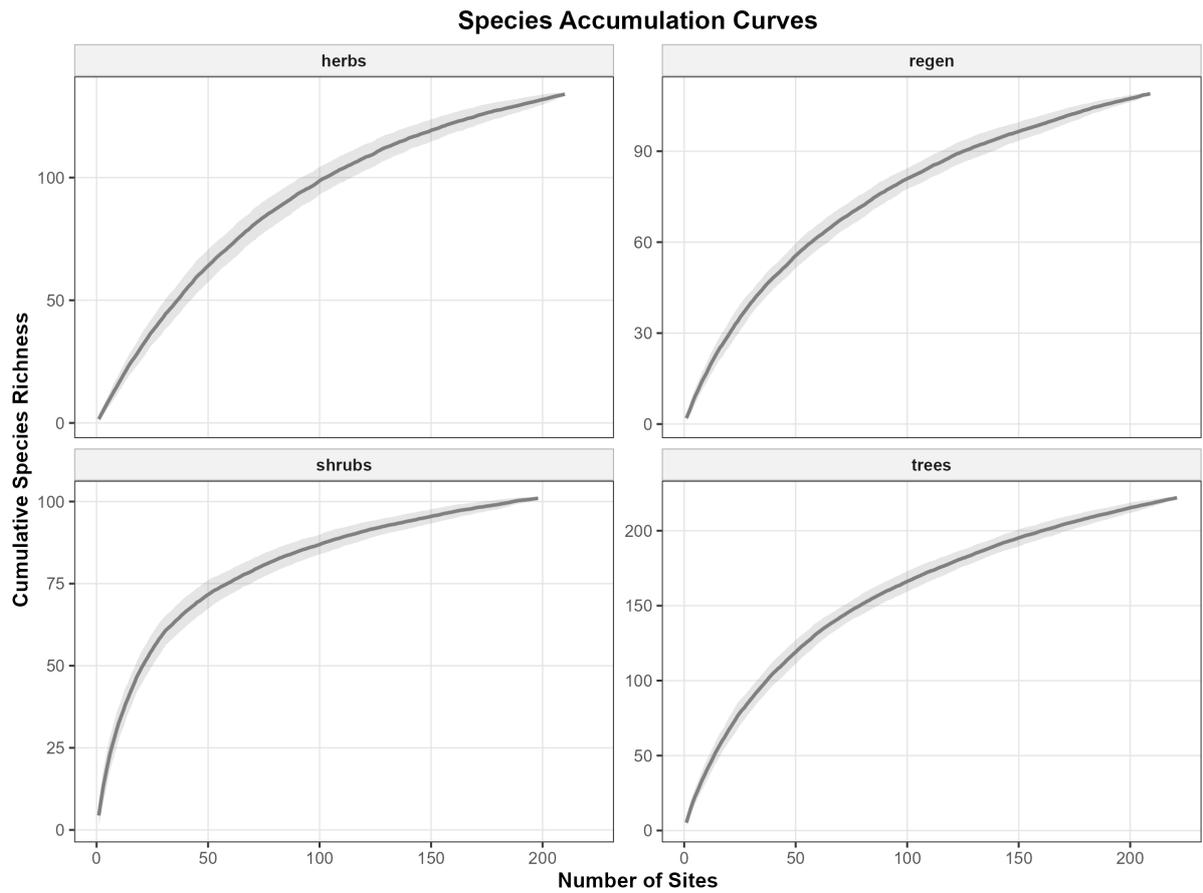


Figure 4: Sample-based species accumulation curves for herbs, regeneration, shrubs and trees. Solid lines indicate mean richness across random permutations; shaded envelopes show 95% confidence intervals.

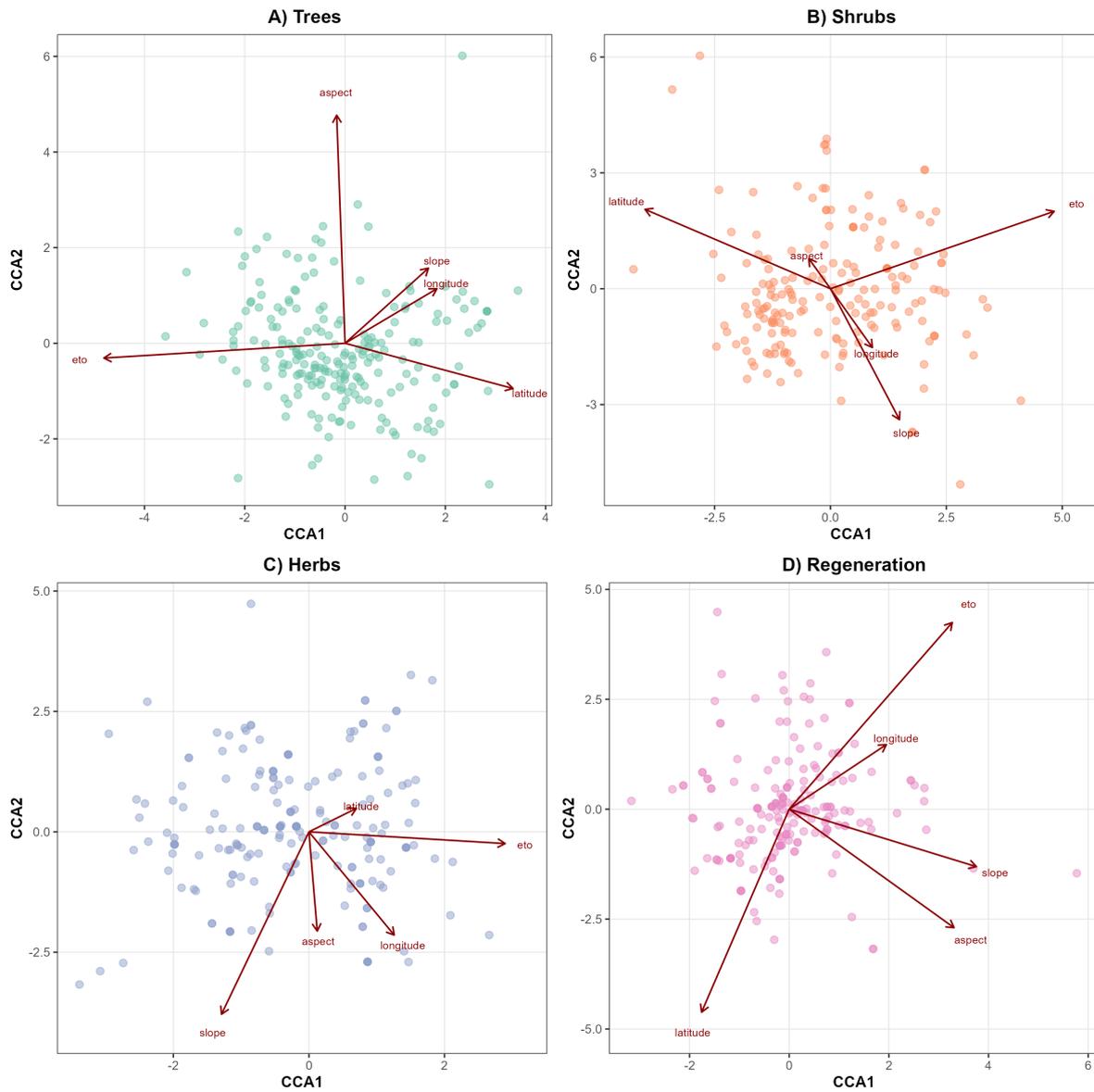


Figure 5: Canonical correspondence analysis (CCA) biplots for (A) trees, (B) shrubs, (C) herbs and (D) regeneration. Points represent plots and arrows denote environmental predictors retained after variance inflation factor screening (threshold = 10). Arrow length indicates the strength of correlations with ordination axes.

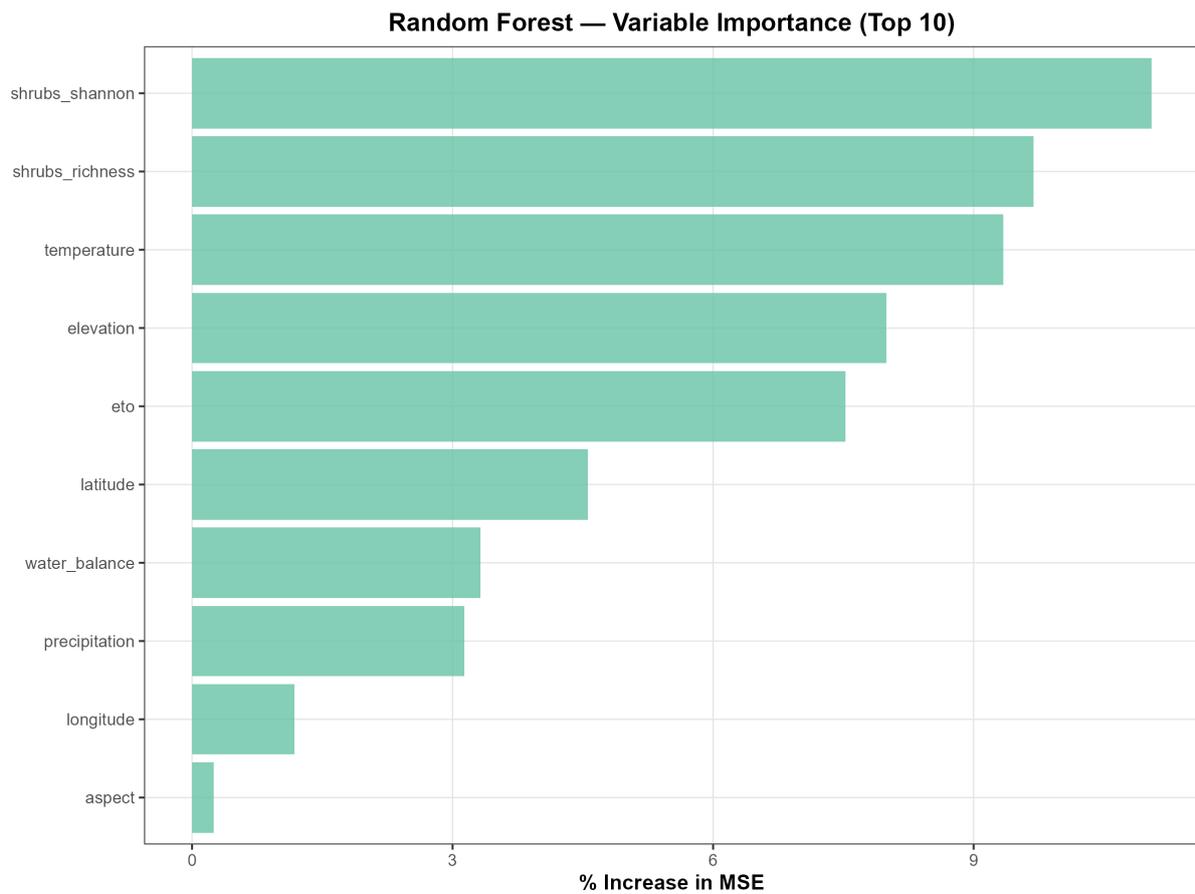


Figure 6: Random-forest variable-importance ranking for predictors of regeneration richness. Bars show percentage increase in mean squared error ($\%IncMSE$) following permutation of each variable during five-fold cross-validation. Higher values indicate stronger influence on predictive accuracy.

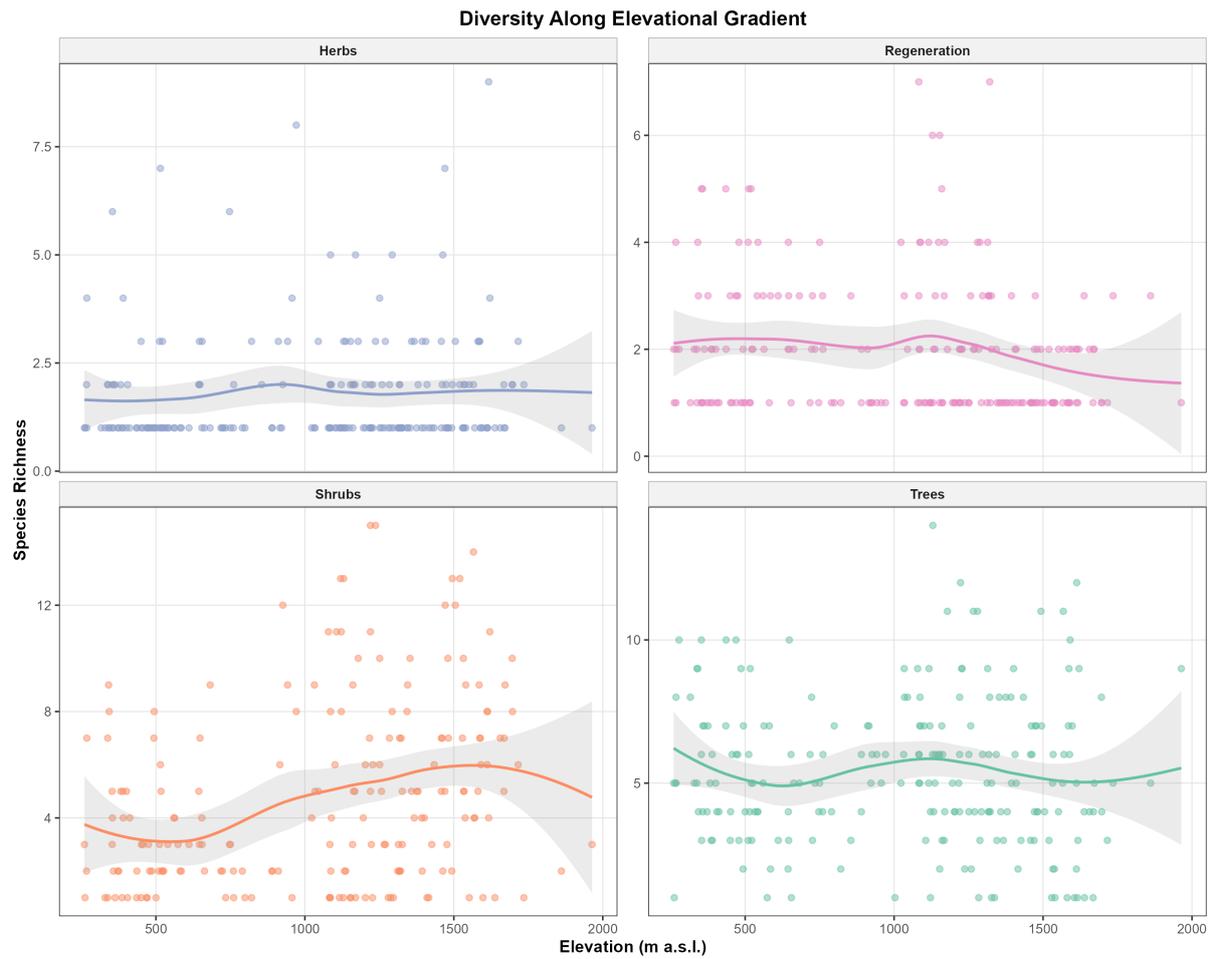


Figure 7: Variation in plot-level species richness along the elevational gradient for herbs, regeneration, shrubs and trees. Curves represent locally weighted regression fits with 95% confidence bands.

472 **Supporting Information**

473 Additional supporting information may be found online in the Supporting Information section:

474 Appendix S1. Variance inflation factor screening by stratum.

475 Appendix S2. Full PERMANOVA and dispersion outputs.

476 Appendix S3. Envfit results for all strata.

477 Appendix S4. Indicator species list.

478 Appendix S5. Machine-learning model parameters.

479 Appendix S6. Variable importance outputs.

480 Appendix S7. Cross-validation fold results.

481 Appendix S8. Diversity correlation matrix.

482 Appendix S9. Beta diversity metrics.

483 Appendix S10. Community matrix diagnostics.