Understanding how biological communities are shaped is a central tenet of community ecology. Recent evidence highlights the potential of decoupling diversity spatial autocorrelation into its positive and negative components to reveal community assembly processes that would otherwise remain undetected, as well as to improve understanding of their impacts on different facets of diversity. Yet, such approaches have only been implemented to investigate the effects of a few assembly drivers on a small number of diversity components. Here, we used high Andean wetland plant communities over a strong latitudinal gradient to investigate the effects of various ecological factors on spatial autocorrelation patterns of nine community metrics with different informative values, including measures of richness, dominance, evenness and beta-diversity. By combining Moran’s Eigenvector Maps, partial least squares structural equation modeling, and regression analyses, we revealed two groups of community parameters presenting contrasting spatial patterns due to specific sensitivities to ecological factors. While environmental variation and wetland connectivity increased positive spatial autocorrelation in richness and dominance-related parameters, species co-occurrence promoted negative spatial autocorrelation in evenness-related parameters. These results offer new insights regarding both how ecological processes affect species assembly, as well as the information captured by classical taxonomic parameters.

Understanding how biological communities are shaped is necessary for biodiversity conservation1–3, safeguarding ecosystem functions and services4, and predicting responses to global change5. A large component of current research in community ecology is thus devoted to disentangling processes involved in community assembly. However, gathering direct empirical evidence of the ecological processes at play is almost impossible under natural conditions6. As a result, they are usually assessed indirectly through pattern-process relationship approaches7,8, which rely on the premise that assembly mechanisms leave particular signatures on communities, and that ecologists are able to unequivocally identify these footprints.

Among the approaches for disentangling mechanisms underpinning community diversity9–12 that have gained popularity in recent years is the investigation of diversity patterns across multiple spatial scales. This strategy rests on the fact that assembly processes generate nonrandom spatial structures, leaving footprints at different spatial scales12–15. Two major groups of ecological processes can be distinguished in this respect: those with broad-scale consequences for species distribution and diversity and those with fine/local scale effects. The former group corresponds to processes generating positive spatial autocorrelations in community diversity, which refers to situations in which diversity levels at nearby locations are more similar than expected by chance16. Such processes include contagious processes16,17, such as dispersal, that generate influences from surrounding sites on community composition, and environmental filtering driven by broad-scale gradients, which similarly leads to commonalities among sites in close geographic proximity due to their high environmental similarity16. The latter group incorporates mechanisms such as biotic interactions with little or no effect beyond the local scale. Because such mechanisms can create singular patterns locally, they are assumed to drive dissimilarities among nearby sites, that is negative spatial autocorrelations13.

Recently, Biswas et al.14 proposed a conceptual framework that capitalizes on these characteristics to identify assembly processes operating at various spatial scales and analyze their spatial pattern effects on different diversity components. The approach relies on the decoupling of diversity spatial autocorrelation into its positive ($S^+ (x)$) and
### Table 1. Expected spatial signatures of community assembly processes on different community characteristics.

S+(x): positive spatial autocorrelation, S−(x): negative spatial autocorrelation.

| Assembly process          | Expected spatial signature | Effects on community characteristics | Most prominent expected pattern                |
|---------------------------|-----------------------------|---------------------------------------|-----------------------------------------------|
| Dispersal                 | Driver of S+(x). Dispersal is a contagious process, generating similarities among nearby communities | Dispersal is expected to increase alpha-diversity in general. The theory of Island Biogeography postulates a major effect of dispersal on richness. Analyses of plant, invertebrate and macroinvertebrate communities suggest a direct effect of migration on species richness. Dispersal is also a main driver of community assembly increasing similarity among connected communities. Empirical studies indicate a major role of dispersal on community uniqueness. | Driver of S+(x) in richness and local contribution to beta-diversity |
| Environmental filtering   | Driver of S+(x) essentially. Environmental filtering can provoke similarities among nearby communities through induced spatial dependence due to the non-random distribution of the environmental variables acting as filters. Such effects are expected to be important along environmental gradients. Occasional driver of S+(x) | According to empirical evidence, environmental filtering more strongly influences species richness than evenness. Environmental heterogeneity is also a common driver of community divergence. A major role of environmental variation on community uniqueness is commonly reported. | Driver of S+(x) in richness and LCBD |
| Species interactions (competition / facilitation) | Potential driver of S+(x). Species interactions is a community assembly driver with no expected effects beyond the local scale. It can thus lead to dissimilarities among nearby communities | Both competition and facilitation alter species number and relative abundance. However, theoretical and empirical evidence indicates that they have a more direct and stronger impact on evenness than on richness. Considering that competition is expected to alter dominance patterns in communities, strong effects can be expected on diversity metrics emphasizing dominance. While species interactions can increase differences among ecological communities, their effects on ecological uniqueness are not established. | Driver of S+(x) in evenness- and dominance-related indices |
| Ecological drift          | Driver of S+(x). Process with no expected effects beyond the local scale. It can thus lead to dissimilarities among nearby communities | Ecological drift provokes random fluctuations in species abundances, lowers diversity within communities and increases differences among ecological communities. The specific sensitivity of richness and abundance-related indices to drift is not well established. | Driver of S+(x) in LCBD and alpha diversity parameters |

negative components (S−(x)). It has the potential to reveal impacts that would remain undetected without such partitioning. For instance, it revealed that environmental disturbance enhances negative spatial autocorrelation in community diversity without necessarily affecting overall spatial autocorrelation. In addition, it showed that disturbance provokes stronger responses in evenness than in richness. These examples demonstrate that the autocorrelation decomposition approach to characterize the impacts of a range of ecological factors on spatial patterns of taxonomic community parameters and identify which they alter the most. To do so, we analyzed plant communities from high Andean wetlands in the Norte Chico region in Chile, which offer an excellent model for the purpose of our study. Since these ecosystems are insular formations distributed across a marked aridity gradient, various processes are expected to leave imprints at a range of spatial scales. These include dispersal restriction and environmental filtering, two potentially important assembly mechanisms with predicted broad-scale imprints (Table 1), and ecological drift and plant-plant interactions with projected influences at the local scale (i.e., individual wetlands, Table 1). The effects of dispersal on S+(x) is suggested by the patchy and highly isolated nature of these habitats, which likely allow migration only between wetlands in relatively close geographic proximity, and may thus precipitate stronger similarities in community composition and diversity among closest sites. The broad-scale impacts of environmental filtering are anticipated due to the pronounced latitudinal climate gradient in the region, which gradually transitions from Mediterranean in the south to arid in the north. The importance of drift, the random changes in species relative abundances, is substantiated by theory and empirical evidence suggesting that it is a major driver of diversity in island-type habitats. Its actual importance may depend on landscape factors influencing population size within communities and thereby risks of demographic stochasticity. As for plant-plant interactions, they are well established key drivers of alpine vegetation. At high elevations, where assemblages are subjected to harsh climatic conditions, positive interactions in particular are expected to assume high importance. They are promoted by facilitator organisms, such as cushion plants, that act as foundation species, stabilizing and improving local abiotic conditions.
To investigate the spatial footprints of these processes on high Andean plant diversity, we analyzed how associated ecological factors (wetland connectivity for dispersal, environmental variation for environmental filtering, wetland size for drift, and species co-occurrence for species interactions influence spatial patterns in community characteristics. Since assembly processes can alter different aspects of the communities (Table 1), we explored ecological factor effects on various alpha-diversity measures (see Supplementary Table S1 online), including measures of richness, dominance (Shannon entropy, Shannon diversity and Simpson diversity), evenness (Piou's evenness, Shannon evenness and Simpson evenness), as well as on a beta-diversity parameter (i.e., local contribution to beta-diversity, LCBD). We postulate that ecological factors associated with broad-scale processes (wetland connectivity and environmental variation) will drive $S^+(x)$ and have stronger effects on richness-related parameters and LCBD (Table 1). In addition, we expect ecological factors associated with fine-scale processes (wetland size and species co-occurrence) to generate $S^-(x)$ in LCBD, evenness and dominance-related parameters (Table 1). To test these predictions, we analyzed positive and negative autocorrelation patterns of each of the community metrics using Moran Eigenvector Maps. Then, we used partial least squares structural equation modeling (PLS-SEM) to estimate the influence of the ecological factors (abiotic environmental variation, species co-occurrence, wetland connectivity, and size) on each plant community parameters. Finally, we investigated whether the influence of ecological factors drove negative or positive spatial autocorrelation in community parameters by investigating the effects of the PLS-SEM path coefficients after excluding potential confounding effects due to the relatedness of the community parameters.

Results
Spatial structure of the diversity parameters. All diversity indices displayed higher levels of $S^+(x)$ than $S^-(x)$ (Fig. 1). Substantial variation was revealed for both autocorrelation components, however, as indicated by coefficients of variation (CVs) of 50% and 38.5% for $S^+$ and $S^-$, respectively. Two groups of alpha-diversity indices stood out based on their spatial patterns. The first, comprising $N_0, N_1, H$, and $N_2$, displayed the highest and only significant levels of $S^+(x)$, but were characterized by the lowest levels of $S^-$, while the second, comprising $E_{10}, E_{20}, J$ and TB, demonstrated the opposite. Autocorrelation patterns of LCBD were intermediate between the two groups of alpha-diversity metrics. Overall, the community metrics displaying the highest $S^+(x)$ tended to be associated with low levels of $S^-(x)$, and vice versa, as indicated by the strong negative relationship between the two autocorrelation components (Pearson correlation: $r = -0.90$, likelihood ratio test of a mixed model of $S^+(x)$ in relation to $S^-(x)$ including the correlation matrix of community metrics as a random effect: $LR = 16.62, df = 1, P < 0.001$). Neither the levels of $S^+(x)$ nor those of $S^-(x)$ were influenced by the covariance of the community metrics (see Supplementary Table S2 online).
Table 2. Influence of wetland connectivity and size, species co-occurrence and environmental variability on each plant community parameter estimated by partial least squares structural equation modeling. The confidence intervals and significance of path coefficients of each ecological factor were evaluated based on 10,000 bootstrapping iterations. Significant path coefficients are in bold. Models were carried out separately for each community metric. N0: Richness; H: Shannon entropy; N1: Shannon diversity; N2: Simpson diversity; E10: Shannon evenness; E20: Simpson evenness; J: Pielou's evenness, TB: Total biomass and LCBD: Local connectivity effects, and the S−(x) levels with effects of species co-occurrence. Wetland connectivity and environmental variation both increased S+(x) of the community indices they positively influenced (Table 3, Figs. 2a,c). Species co-occurrence effects were positive for richness and dominance-influenced parameters (i.e., H and N1), but negative for evenness metrics (E10 and E20). Overall, wetland connectivity and environmental variation had the strongest effects on the community parameters with mean absolute effects being at least three times higher than those of species co-occurrence and wetland size (e.g., mean of path coefficient absolute values in Table 2).

### Influence of local ecological factors on high Andean plant community characteristics.

Among the four ecological factors, only wetland connectivity and size had consistent effects on diversity indices (Table 2). Connectivity increased all community metrics, and significantly so in the case of N1 and N0. In contrast, the estimated effects of wetland size were negative, indicating that increases in wetland size tended to produce a decrease in the community parameters and significantly so in N0. The impacts of environmental variation and species co-occurrence were inconsistent, with path coefficients ranging from negative to positive and characterized by very large CVs (i.e., 243 and 833% respectively, Table 2). Environmental effects were negative except for N0, N1 and N2 (Table 2, Fig. 2b). Species co-occurrence effects were positive for richness and dominance-influenced parameters (i.e., H and N1), but negative for evenness metrics (E10 and E20). Overall, wetland connectivity and environmental variation had the strongest effects on the community parameters with mean absolute effects being at least three times higher than those of species co-occurrence and wetland size (e.g., mean of path coefficient absolute values in Table 2).

### Effects of local ecological factors on positive and negative spatial autocorrelation components of diversity indices.

Three of the four ecological factors increased spatial autocorrelation. Significant relationships between the ecological factor effects (i.e., PLS-SEM path coefficients) and levels of autocorrelation of the community parameters were actually revealed for wetland connectivity, environmental variation and species co-occurrence (Table 3). S+(x) levels showed a significant relationship with environmental variation and wetland connectivity effects, and the S−(x) levels with effects of species co-occurrence. Wetland connectivity and environmental variation both increased S+(x) of the community indices that they positively influenced (Table 3, Fig. 2a,b), and particularly N0, N1, N2 and H (Fig. 2a,b,d). S−(x) levels increased in response to negative effects of species occurrence on community indices, leading to increased levels of S+(x) in E10, E20 and J (Fig. 2a,c).

### Discussion

By investigating the effects of ecological determinants on the spatial autocorrelation components of different diversity indices, we were able to reveal spatial footprints of assembly drivers and identify groups of diversity metrics presenting specific responses. These results thus provide new insights both on how ecological processes affect species assembly as well as the information captured by classical taxonomic diversity indices.

Here, we identified a range of drivers of S+(x) and S−(x) in high Andean plant communities. Environmental variation and wetland connectivity were two factors anticipated to promote similarity in community diversity among wetlands in close geographic proximity36,38. The fact that these factors had the strongest detected effects on community parameters is consistent with previous evidence that suggests they each play a key role in spatial patterns of high Andean wetland communities and diversity7,19,20,39,40. Their impacts on S+(x) in community metrics is also in line with literature associating dispersal restriction and environmental filtering with broad spatial scales4,14,11, when dispersal is a limiting factor in structuring communities42 and in the presence of environmental gradients46, as was the case here. Our results demonstrate that not all diversity components were equally influenced by these factors, however. Environmental variation and wetland connectivity increased similarity among wetlands in close geographic proximity in species richness and dominance-related indices (N0, N2, N1 and H), but not in evenness. This adds to the mounting body of evidence showing that richness and evenness respond differently to a wide range of factors and mechanisms4,14,37,41,43–39. As predicted by theoretical literature and earlier empirical results4,41,43,47,48, we found that habitat connectivity increased species richness and S+(x) levels in...
this diversity component. High levels of connectivity also reduced community dominance (i.e., in $N_1$), which is consistent with theoretical expectations that dominance increases with low migration rates, and vice versa.

A noteworthy finding was that environmental effects promoted different spatial patterns on the community parameters depending on their specific impacts. They increased $S^+(x)$ in the community metrics that they positively influenced (i.e., $N_0$, $N_1$, $N_2$ and $H$). $S^+(x)$ and $S^-(x)$ refer to positive and negative spatial autocorrelations, respectively. $N_0$: Richness; $H$: Shannon entropy; $N_1$: Shannon diversity; $N_2$: Simpson diversity; $E_{10}$: Shannon evenness; $E_{20}$: Simpson evenness; $J$: Pielou’s evenness, TB: Total biomass and LCBD: Local Contribution to Beta-Diversity.

**Figure 2.** Relationships between the effects of ecological factors on community characteristics and the levels of spatial autocorrelation displayed by the latter. (a, b, c) Biplots for significant relationships of the autocorrelation levels of the community metrics by the amplitude of ecological factor’s path coefficients. Dots indicate the position of each community parameter. (d) Representation of the ecological factor effects. Bar sizes are proportional to detected effects. The colors indicate whether increases in autocorrelation are mediated by positive or negative effects of ecological factors on diversity indices. Positive effects are shown in red, negative in blue, and null in white. For instance, both wetland connectivity and environmental variation increased $S^+(x)$ of diversity indices that they positively influenced (i.e., $N_0$, $N_1$, $N_2$ and $H$). $S^+(x)$ and $S^-(x)$ refer to positive and negative spatial autocorrelations, respectively. $N_0$: Richness; $H$: Shannon entropy; $N_1$: Shannon diversity; $N_2$: Simpson diversity; $E_{10}$: Shannon evenness; $E_{20}$: Simpson evenness; $J$: Pielou’s evenness, TB: Total biomass and LCBD: Local Contribution to Beta-Diversity.
occurrence on S-(x), they were found to promote dissimilarities among geographically close wetlands in evenness suggests that it reflects the development of competitively superior species. Regarding the effects of species co-

Table 3. Effects of ecological factors on autocorrelation levels of plant community parameters according to an exhaustive model selection approach based on the Akaike information criterion corrected for small sample size. Ecological factor effects were tested separately for the positive (S’(x)) and negative (S’(x)) spatial autocorrelation components of the community parameters. The models used the autocorrelation levels of each community parameter as the dependent variable and the PLS-SEM path coefficients measuring the direct effects of the ecological factors on each community characteristic as predictors.

| Spatial autocorrelation component | Adjusted R² | F  | dfnum,den | P   | Ecological factor | β ± sd | t     | P     |
|----------------------------------|-------------|----|-----------|-----|------------------|--------|-------|-------|
| S’(x)                            | 0.87        | 27.49 | 2.6       | < 0.001 | Wetland connectivity | 0.064 ± 0.017 | 3.757 | 0.009 |
| S’(x)                            | 0.63        | 14.34 | 1.7       | 0.006    | Species co-occurrence | 0.028 ± 0.007 | 3.786 | 0.007 |

A striking finding of our study is the clear separation of two groups of alpha-diversity parameters presenting contrasting spatial patterns due to their specific sensitivities to ecological factors. Since our analyses discarded any confounding effects of the non-independency of the indices, this result cannot be considered as a simple mathematical artefact of their relationships, and thus demonstrates that assembly mechanisms tend to influence specific aspects of diversity. Understanding the informative value of diversity metrics is a longstanding issue in ecology60–67, and debate is ongoing regarding which measures are optimal65,67–69, as well as whether a single index could be universally applied for comparative purposes69,70. The fact that two groups of diversity metrics bore different spatial footprints demonstrates that each group is sensitive to different assembly processes, and encourages their combined use to aid in a better and more thorough understanding of the ecological processes that sustain community assembly. In addition, it shows that each diversity index holds specific potential for conservation and restoration purposes by motivating for either local or landscape-scale actions. While emerging trends from this and previous studies suggest that evenness tends to be particularly sensitive to local/fine-scale processes and richness to broader scale mechanisms, our understanding of how assembly factors generate and alter spatial patterns of different diversity facets is still limited to a few studies61,63,18,71. More empirical evidence from a wider range of ecosystems, including non-patchy habitats, is therefore required to establish the generality of these patterns. This will be essential to determining the specific potential of diversity parameters to decipher processes shaping communities.

Methods

Data collection and production. Sampling and processing of high Andean wetland plant communities. We used plant data of Bertin et al.20 and Pfeiffer et al.40 collected in 21 high Andean wetlands along a 600 km latitudinal range (between 26°S and 32°S) in north-central Chile between altitudes of 2852 and 4307 m a.s.l. (See Fig. 2 in Bertin et al.20). High Andean wetlands are permanent features formed by groundwater up-welling and occur in the low Alpine and sub-Alpine belts of the Central Andes72, fed by glacial streams, snow-melt, and rainfall72. These insular features contrast markedly from the arid grassland matrix in which they are found73. In the Norte Chico region of Chile, the location of this study, they occur along a latitudinal climatic gradient that varies from hyper-arid in the north to Mediterranean in the south, with mean annual precipitation extremes ranging from 35 to 200 mm, respectively. They support a wide variety plants and animals, including a number of endemic and rare species22,74.

Plant assemblages were characterized for each wetland from five 30 × 30 cm vegetation quadrats20. The biomass (g/m²) of each species was estimated by summing dried biomass of all individuals present within a quadrat after complete drying of the plant samples at 70 °C. A total of 52 species belonging to 21 families were identified. The plant database can be found as Supplementary Table S3 online.

Community parameters. Nine community metrics (see Supplementary Table S1 online) were estimated after pooling data from the five vegetation quadrats. They included total biomass estimated as biomass (dry weight in gr/m2) of all species present in the community samples, richness, three dominance-related metrics (Shannon entropy, Shannon diversity; Simpson diversity) and three evenness-related metrics (Shannon evenness, Simpson
evenness, Pielou’s evenness) and the local contribution to beta-diversity calculated from the Hellinger transformed community data using the function beta.div in the adespatial package89.

**Ecological variables.** We assessed characteristics likely linked to environmental filtering, dispersal rates, drift, and species interactions. For environmental filtering effects, we used a subset of landform and climate related variables derived from the original datasets used by Bertin et al.19 and Pfeiffer et al.40, and which can be found in Supplementary Table S5 online. Only non-redundant variables with apparent influence on the diversity parameters were considered. These were identified by performing partial least square (PLS) regressions on the different diversity metrics. Only environmental variables that contributed strongly to the retained PLS components (defined using a leave-one out cross-validation procedure and by identifying the first local minimum of the root-mean-squared error29) were then retained for subsequent analyses. They included altitude, slope, mean annual wind speed (MAWS), mean annual number of snow days (SnowNDays), mean productivity (NDVI), and productivity variation (δNDVI).

We used graph-based connectivity metrics as variables related to dispersal rates. These were quantified for each of the wetlands in this study using GRAPHAB 1.076, by considering all other wetlands occurring within a 20 km radius from the focal wetland according to the wetland map generated by Bertin et al.30. The connectivity measures calculated were node degree (Dg, the number of patches close to the focal patch77), closeness centrality (CCe, the mean distance from the focal patch to all other patches of its component79), and betweenness centrality (BC, the sum of the shortest paths through the focal patch68,80). We used wetland size and species co-occurrence as proxies for drift risk and plant species interactions, respectively. Wetland size was assessed based on remotely-sensed NDVI derived from LandSat 8 OLI satellite imagery (see Bertin et al.30). Species co-occurrences were analyzed by calculating an average C-score for each wetland from the five plots. The C-score metric measures the level of segregation of all species pairs, which increases as average co-occurrences between all of the species pairs in the matrix decrease81. Thus, to allow an intuitive interpretation, the effects that we report for species co-occurrence are the additive opposite of the C-score effects. The C-score values were calculated with the R-package EcoSimR81.

**Statistical analyses.** Spatial structure analyses of the plant community parameters. We estimated spatial autocorrelation components of the community parameters using Moran’s Eigenvector Maps88 as implemented in the R packages spdep82 and adespatial83. This framework allows producing spatial predictors, the MEMs, which model autocorrelation patterns at different spatial scales. They are obtained by eigen-decomposition of a spatial weighting matrix coding for connections between sites (i.e. wetlands in this instance), which produces orthogonal eigenvectors maximizing the spatial autocorrelation computed by Moran’s I84. When associated with positive eigenvalues, the MEMs represent positive autocorrelation and vice versa13. The spatial weighting matrix that we used to generate the MEMs considered connections among all wetlands for which strength was inversely proportional to their Euclidean distance. To estimate S+(x) and S−(x) of community indices, we followed the procedure described in Biswas et al.14,15, which involved summing the squared correlations between each diversity parameter and the positive and negative MEMs weighted by their associated eigenvalue.

**Influence of ecological factors on plant community parameters.** We investigated the influence of each ecological factor (wetland connectivity and size, species co-occurrence, and environmental variability) on community indices by carrying out PLS-SEM85. This is a comprehensive technique based on the use of a variable covariance matrix that identifies relationships and causality between variables86 with minimum requirements regarding measurement scales, sample sizes, and residual distributions87. The measurement model (i.e. the outer model, see Supplementary Figure S1 online) served to define latent variables representing the four ecological factors (wetland size, wetland connectivity, environmental variability, species co-occurrence) from the 12 measured ecological variables85. Environmental variability, wetland size, and connectivity were defined according to a formative mode and species co-occurrence using a reflective mode (see Supplementary Figure S1 online). Inspections of the loadings of the ecological variables on the latent variables indicated that the latent variables for wetland size and connectivity increased as wetland size and connectivity increased, respectively. The inner or structural model was constructed to estimate the effects of each of the four ecological factors on plant community characteristics (see Supplementary Figure S1 online). Eight separate PLS-SEM models were run to investigate these effects on each of the community metrics. Confidence intervals and significance of path coefficients were evaluated based on 10,000 bootstrapping iterations. These analyses were performed using the semPLS R-packages85.

**Effects of assembly driving factors on positive and negative spatial autocorrelation components of community indices.** To investigate whether the assembly driving factors influenced autocorrelation patterns of the community indices, we tested the relationship between the effects of the ecological factors (as estimated by their path coefficients) and the autocorrelation levels of the diversity parameters. To account for relatedness between the community parameters, we first analyzed the effects of their covariation on S+(x) and S−(x) levels. The pairwise Pearson’s correlation matrix of the plant indices was transformed through principal coordinate decomposition (PCoA16,89) using the ecodist R package89. Then, we analyzed the effects of the PCoA eigenvectors so-produced on S+(x) and S−(x) separately by carrying out generalized linear models using the R leaps package90. In these analyses, the autocorrelation levels of each community parameter were used as the dependent variables and the eigenvectors as the predictors. We ran the regsubset function, which performs an automatic selection procedure to look for the subset of eigenvectors that best explained the autocorrelation variation of the community indices. The best-fitting model, identified through a performance analysis performed with the AICcmodavg package85 and using the Akaike’s Information Criterion corrected for small sample size (AICc), was then tested for sig-
nificance. Since these analyses did not reveal any impact of the relatedness between the community indices on their autocorrelation levels (see Supplementary Table S2 online), influences of the plant community parameters on $S^+$ and $S^-$ were analyzed through multiple linear regressions. The dependent variable of these analyses were the autocorrelation levels of each community parameter and the predictors the PLS-SEM path coefficients measuring the direct effects of the ecological factors on each community characteristic. For these analyses, we also performed an exhaustive model selection to search for the subset of ecological factors that best explained the autocorrelation levels using, as above, the regsubset function and based on the AICc criteria. Normality of the residuals was verified using Shapiro tests.

Data availability

All the data used in the analyses are provided in Supplementary Tables S3, S4 and S5 of the Supplementary Information file.

Received: 8 February 2022; Accepted: 5 August 2022

Published online: 12 August 2022

References

1. Rudnick, D. A. et al. The role of landscape connectivity in planning and implementing conservation and restoration priorities. *Issues Ecol.* 16, 1–23 (2012).
2. Brudvig, L. A. Interpreting the effects of landscape connectivity on community diversity. *J. Veg. Sci.* 27, 4–5 (2016).
3. Socolar, J. R.; Gilroy, J. J.; Kunin, W. E.; & Edwards, D. P. How should beta-diversity inform biodiversity conservation? *Trends Ecol. Evol.* 31, 67–80 (2016).
4. Leibold, M. A., Chase, J. M.; & Ernst, S. K. M. Community assembly and the functioning of ecosystems: how metacommunity dynamics influence the direct effects of the ecological factors on each community characteristic. For these analyses, we also performed an exhaustive model selection to search for the subset of ecological factors that best explained the autocorrelation levels using, as above, the regsubset function and based on the AICc criteria. Normality of the residuals was verified using Shapiro tests.

17. Legendre, P.; & De Cáceres, M. Beta diversity as the variance of community data: Dissimilarity coefficients and partitioning. *Ecography* 35, 517–537 (2012).
21. Souvignet, M., Oyarzún, R., Verbist, K. M. J.; Gaese, H. & Heinrich, J. Hydro-meteorological trends in semi-arid north-central Chile (29–32°S): Water resources implications for a fragile Andean region. *Proc. Biol. Sci.* 288, 1–10 (2011).
23. Zúñiga, B.; Storic, B.; Gutiérrez, J. R.; & López, D. Climatic characteristics of the semi-arid Coquimbo Region in Chile. *J. Arid Environ.* 96, 913–924 (2015).
26. Anthelme, F.; & Dangles, O. Plant–plant interactions in tropical alpine environments. *Perspect. Plant Ecol.* 28, 1–10 (2017).
34. Seabloom, E. W.; BJørnstad, O. N.; Bolker, B. M.; & Reichman, O. J. Spatial signature of environmental heterogeneity, dispersal, and competition in successional grasslands. *Ecol. Monogr.* 75, 199–214 (2005).
35. Seabloom, E. W.; BJørnstad, O. N.; Bolker, B. M.; & Reichman, O. J. Spatial signature of environmental heterogeneity, dispersal, and competition in successional grasslands. *Ecol. Monogr.* 75, 199–214 (2005).

37. Wilsey, B. & Stirling, G. Species richness and evenness respond in a different manner to propagule density in developing prairie microcosm communities. Plant Ecol. 190, 259–273 (2007).
38. Schamp, B. S., Arnott, S. E. & Joslin, K. L. Dispersal strength influences zooplankton co-occurrence patterns in experimental mesocosms. Ecology 96, 1074–1083 (2015).
39. Troncoso, A. J., Bertin, A., Osorio, R., Arancio, G. & Gouin, N. Comparative population genetics of two dominant plant species of high Andean wetlands reveals complex evolutionary histories and conservation perspectives in Chile’s Norte Chico. Conserv. Genet. 18, 1047–1060 (2017).
40. Pfeiffer, V. W. et al. Partitioning genetic and species diversity refines our understanding of species–genetic diversity relationships. Ecol. Evol. 8, 12351–12364 (2018).
41. Bello, F. D. et al. Hierarchical effects of environmental filters on the functional structure of plant communities: A case study in the French Alps. Ecography 36, 393–402 (2013).
42. Moritz, C. et al. Disentangling the role of connectivity, environmental filtering, and spatial structure on metacommunity dynamics. Oikos 122, 1401–1410 (2013).
43. Wilsey, B. J. & Potvin, C. Biodiversity and ecosystem functioning: Importance of species evenness in an old field. Ecology 81, 887–892 (2000).
44. Stirling, G. & Wilsey, B. Empirical relationships between species richness, evenness, and proportional diversity. Am. Nat. 158, 286–299 (2001).
45. Stevens, R. D. & Willig, M. R. Geographical ecology at the community level: Perspectives on the diversity of new world bats. Ecology 83, 545–560 (2002).
46. Wilsey, B. J. & Polley, H. W. Effects of seed additions and grazing history on diversity and productivity of subhumid grasslands. Ecology 84, 920–931 (2003).
47. Ma, M. Species richness vs evenness: Independent relationship and different responses to edaphic factors. Oikos 111, 192–198 (2005).
48. Schmitz, O. J. Effects of predator hunting mode on grassland ecosystem function. Science 319, 952–954 (2008).
49. Stomp, M., Huissman, J., Mittelbach, G. G., Litchman, E. & Klausmeier, C. A. Large-scale biodiversity patterns in freshwater phytoplankton. Ecology 92, 2096–2107 (2011).
50. Zhang, H. et al. The relationship between species richness and evenness in plant communities along a successional gradient: A study from sub-alpine meadows of the eastern Qinghai-Tibetan plateau, China. PLoS ONE 7, e49024 (2012).
51. Hubbell, S. P. The Unified Neutral Theory of Biodiversity and Biogeography (Princeton University Press, 2001).
52. Young, K. R. in Climate Change and Biodiversity in the Tropical Andes (eds Herzog, S. K., Martínez, R., Jørgensen, P. M. & Tieszen, H.) Ch. 8, 128–140 (Inter-American Institute for Global Change Research, 2011).
53. López-Angüalo, J. et al. Determinants of high mountain plant diversity in the Chilean Andes: From regional to local spatial scales. PLoS ONE 13, e0200216 (2018).
54. MacArthur, R. H. & Wilson, E. O. The Theory of Island Biogeography (Princeton University Press, 1967).
55. Hanski, I. Metapopulation Ecology (Oxford University Press, 1999).
56. Blanchet, F. G., Cazelles, K. & Gravel, D. Co-occurrence is not evidence of ecological interactions. Ecol. Lett. 23, 1050–1063 (2020).
57. Kunte, K. Competition and species diversity: Removal of dominant species increases diversity in Costa Rican butterfly communities. Oikos 117, 69–76 (2008).
58. Dray, S., Legendre, P. & Peres-Neto, P. R. Spatial modelling: A comprehensive framework for principal coordinate analysis of neighbour matrices (PCNM). Ecol. Model. 196, 483–493 (2006).
59. Kikvidze, Z. et al. Linking patterns and processes in alpine plant communities: A global study. Ecology 86, 1395–1400 (2005).
60. Hill, M. O. Diversity and evenness: A unifying notation and its consequences. Ecology 54, 427–432 (1973).
61. Heip, C. H. R., Herman, P. M. J. & Soetart, K. Indices of diversity and evenness. Oceanis 4, 61–87 (1998).
62. Jost, L. Entropy and diversity. Oikos 113, 363–375 (2006).
63. Jost, L. Partitioning diversity into independent alpha and beta components. Ecology 88, 2427–2439 (2007).
64. Jost, L. The relation between evenness and diversity. Diversity 2, 207–232 (2010).
65. Pallmann, P. et al. Assessing group differences in biodiversity by simultaneously testing a user-defined selection of diversity indices. Mol. Ecol. Resour. 12, 1068–1078 (2012).
66. Chao, A. et al. Rarefaction and extrapolation with Hill numbers: A framework for sampling and estimation in species diversity studies. Ecol. Monogr. 84, 45–67 (2014).
67. Harris, W. A. et al. Choosing and using diversity indices: Insights for ecological applications from the german biodiversity exploratories. Ecol. Evol. 4, 3514–3524 (2014).
68. Beisel, J.-N., Usseglio-Polatera, P., Bachmann, V. & Moretou, J.-C. A comparative analysis of evenness index sensitivity. Int. Rev. Hydrobiol. 88, 3–15 (2003).
69. Fedor, P. & Zvaríková, M. in Encyclopedia of Ecology (ed Brian Fath) 337–346 (2019).
70. Gatti, R. C., Amoroso, N. & Monaco, A. Estimating and comparing biodiversity with a single universal metric. Ecol. Model. 424, 8 (2020).
71. Lin, L., Deng, W., Huang, X. & Kang, B. Fish taxonomic, functional, and phylogenetic diversity and their vulnerabilities in the largest river in southeastern China. Ecol. Evol. 11, 11533–11548 (2021).
72. Squeo, F. A., Warner, B. G., Aravena, R. & Espinoza, D. Bofedales: High altitude peatlands of the central Andes. Rev. Chil. Hist. Nat. 79, 245–255 (2006).
73. Villagrán-Mella, R., Aguayo, M., Parra, L. E. & González, A. Relación entre características del hábitat y estructura del ensamble de insectos en humedales palustres urbanos del centro-sur de Chile. Rev. Chil. Hist. Nat. 79, 195–211 (2006).
74. Coronel, J. S., Delecras, S., Maldonado, M., Ollevier, F. & Brendonck, L. Temporary shallow pools in high-Andes ‘bofedal’ peatlands. Arch. Soc. Sci. 57, 85–96 (2004).
75. Wakeling, J. N. & Morris, J. J. A test of significance for partial least squares regression. J. Chemom. 7, 291–304 (1993).
76. Foltète, J.-C., Cazelles, K. & Vuidel, G. A software tool dedicated to the modelling of landscape networks. Environ. Modell. Sofw. 38, 316–327 (2012).
77. Ricotta, C., Stancioli, A., Avena, G. C. & Blasi, C. Quantifying the network connectivity of landscape mosaics: A graph-theoretical approach. Community Ecol. 1, 89–94 (2000).
78. Freeman, L. C. Centrality in social networks conceptual clarification. Soc. Netw. 1, 215–239 (1979).
79. Urban, D. et al. Landscape connectivity: A graph-theoretic perspective. Ecology 82, 1205–1218 (2001).
80. Bodin, O. & Saura, S. Ranking individual habitat patches as connectivity providers: Integrating network analysis and patch removal experiments. Ecol. Model. 221, 2393–2405 (2010).
81. Gottelli, N. J., Hart, E. M. & Ellison, A. M. EcoSimR: Null model analysis for ecological data. R package version 0.1.0. (R Foundation for Statistical Computing, 2015).
82. Bivand, R. S. & Wong, D. W. S. Comparing implementations of global and local indicators of spatial association. TEST 27, 716–748 (2018).
83. Dray, S. et al. adespatial: multivariate multiscale spatial analysis. R package version 0.3-8. (R Foundation for Statistical Computing, 2020).
Acknowledgements

We thank Vincent Diego for advice to analyze covariance effects, A. Baumel for helpful comments on a previous version of the manuscript, and C. Weideman for revising the English of the manuscript. This work was financed by the Dirección de Investigación y Desarrollo de la Universidad de La Serena (DIDULS), DIDULS regular PR192126. A.L. received a scholarship Program/ DOCTORADO BECAS CHILE/2019 - 21190608 from the National Agency for Research and Development (ANID).

Author contributions

A.B. and A.L. conceived this study. A.L. conducted the statistical analyses. A.B. and A.L. drafted the first version of the manuscript, provided intellectual input, and contributed to the final version of the manuscript.

Competing interests

The authors declare no competing interests.

Additional information

Supplementary Information The online version contains supplementary material available at https://doi.org/10.1038/s41598-022-18132-9.

Correspondence and requests for materials should be addressed to A.B.

Reprints and permissions information is available at www.nature.com/reprints.

Publisher’s note Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

Open Access This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article’s Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article’s Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit http://creativecommons.org/licenses/by/4.0/.

© The Author(s) 2022

84. Wagner, H. H. & Dray, S. Generating spatially constrained null models for irregularly spaced data using Moran spectral randomization methods. *Methods Ecol. Evol.* 6, 1169–1178 (2015).
85. Monecke, A. & Leisch, F. semPLS: Structural equation modeling using partial least squares. *J. Stat. Softw.* 48, 1–32 (2012).
86. Zhao, X., Li, Y., Song, H., Jia, Y. & Liu, J. Agents affecting the productivity of pine plantations on the Loess Plateau in China: A study based on structural equation modeling. *Forests* 11, 1328 (2020).
87. Tenenhaus, M., Vinzi, V. E., Chatelin, Y.-M. & Lauro, C. PLS path modeling. *Comput. Stat. Data Anal.* 48, 159–205 (2005).
88. Gower, J. C. & Legendre, P. Metric and euclidean properties of dissimilarity coefficients. *J. Classif.* 3, 3–48 (1986).
89. Gosle, S. C. & Urban, D. L. The ecodist package for dissimilarity-based analysis of ecological data. *J. Stat. Softw.* 22, 1–19 (2007).
90. Lumley, T. & Miller, A. leaps: Regression subset selection. R package version 2.7. http://CRAN.R-project.org/package=leaps (2004).
91. AICcmodavg: Model Selection and Multimodel Inference Based on (Q)AIC(c). R package version 2.3-1. (2019).
92. Freestone, A. L. & Inouye, B. D. Dispersal limitation and environmental heterogeneity shape scale-dependent diversity patterns in plant communities. *Ecology* 87, 2425–2432 (2006).
93. Li, F., Tonkin, J. D. & Haase, P. Local contribution to beta diversity is negatively linked with community-wide dispersal capacity in stream invertebrate communities. *Ecol. Indic.* 108, 105715 (2020).
94. Vilmi, A., Karjalainen, S. M. & Heino, J. Ecological uniqueness of stream and lake diatom communities shows different macro-ecological patterns. *Divers. Distrib.* 23, 1042–1053 (2017).
95. Baldeck, C. A., Tupayachi, R., Sinca, F., Jaramillo, N. J. E. & Asner, G. P. Environmental drivers of tree community turnover in western Amazonian forests. *Ecography* 39, 1089–1099 (2016).
96. Chase, J. M. Stochastic community assembly causes higher biodiversity in more productive environments. *Science* 328, 1388–1391 (2010).
97. Chase, J. M. & Myers, J. A. Entanglement of the importance of ecological niches from stochastic processes across scales. *Philos. Trans. R. Soc. B* 366, 2351–2363 (2011).
98. Segre, H. et al. Competitive exclusion, beta diversity, and deterministic vs. stochastic drivers of community assembly. *Ecol. Lett.* 17, 1400–1408 (2014).
99. Ceschin, F., Bini, L. M. & Padial, A. A. Correlates of fish and aquatic macrophyte beta diversity in the Upper Paraná River floodplain. *Hydrobiologia* 805, 377–389 (2018).
100. Heino, J. et al. Unraveling the correlates of species richness and ecological uniqueness in a metacommunity of urban pond insects. *Ecol. Indic.* 73, 422–431 (2017).
101. Leão, H., Siqueira, T., Torres, N. R. & Montag, L. F. D. A. Ecological uniqueness of fish communities from streams in modified landscapes of Eastern Amazonia. *Ecol. Indic.* 111, 106039 (2020).
102. Vega-Álvarez, J., García-Rodríguez, J. A. & Cayuela, L. Facilitation beyond species richness. *J. Ecol.* 107, 722–734 (2019).