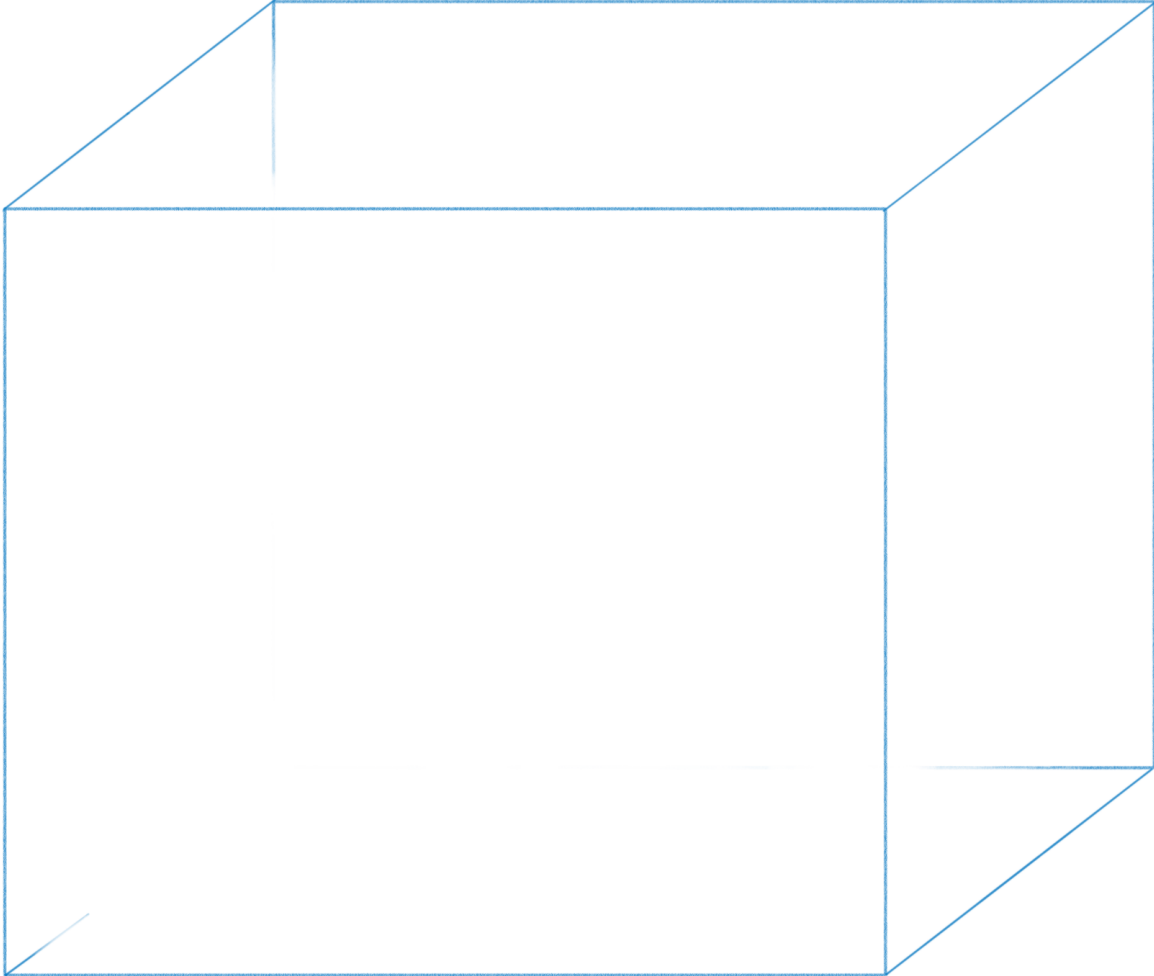


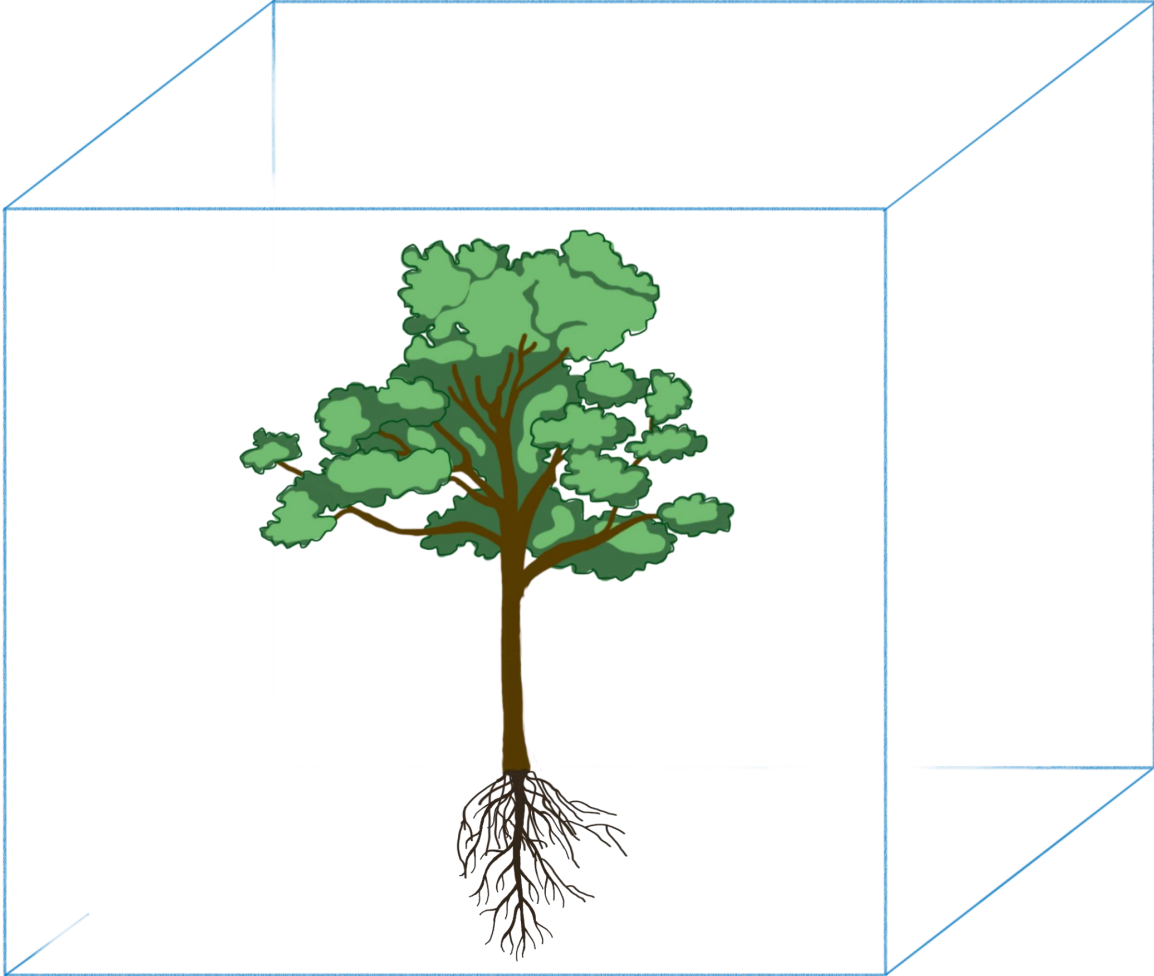
Comparison of the global distribution of functional and phylogenetic diversity in plant communities.

A study to highlight commonalities and differences in the distribution of vascular plants.

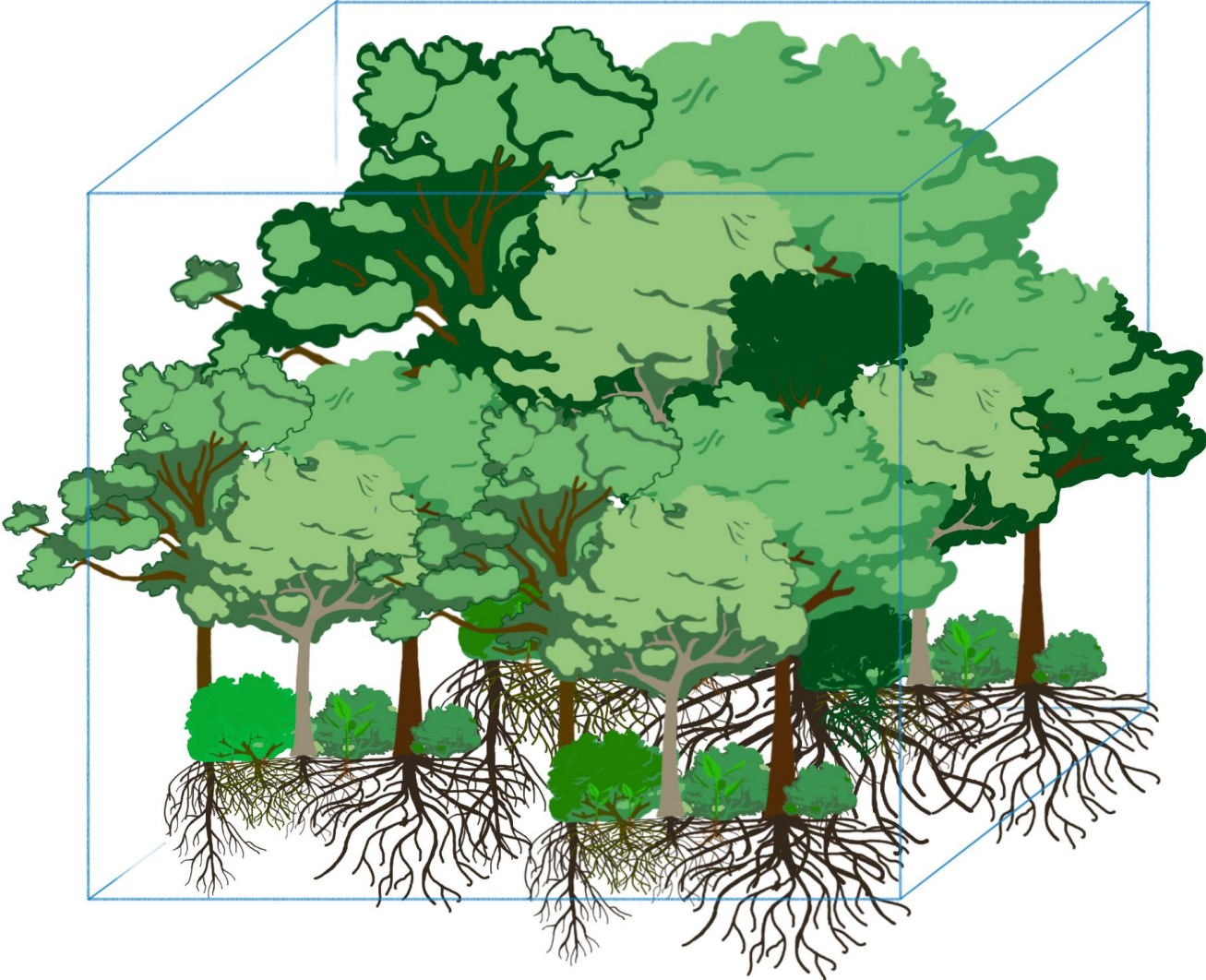
Introduction – Environmental space



Introduction – Environmental space

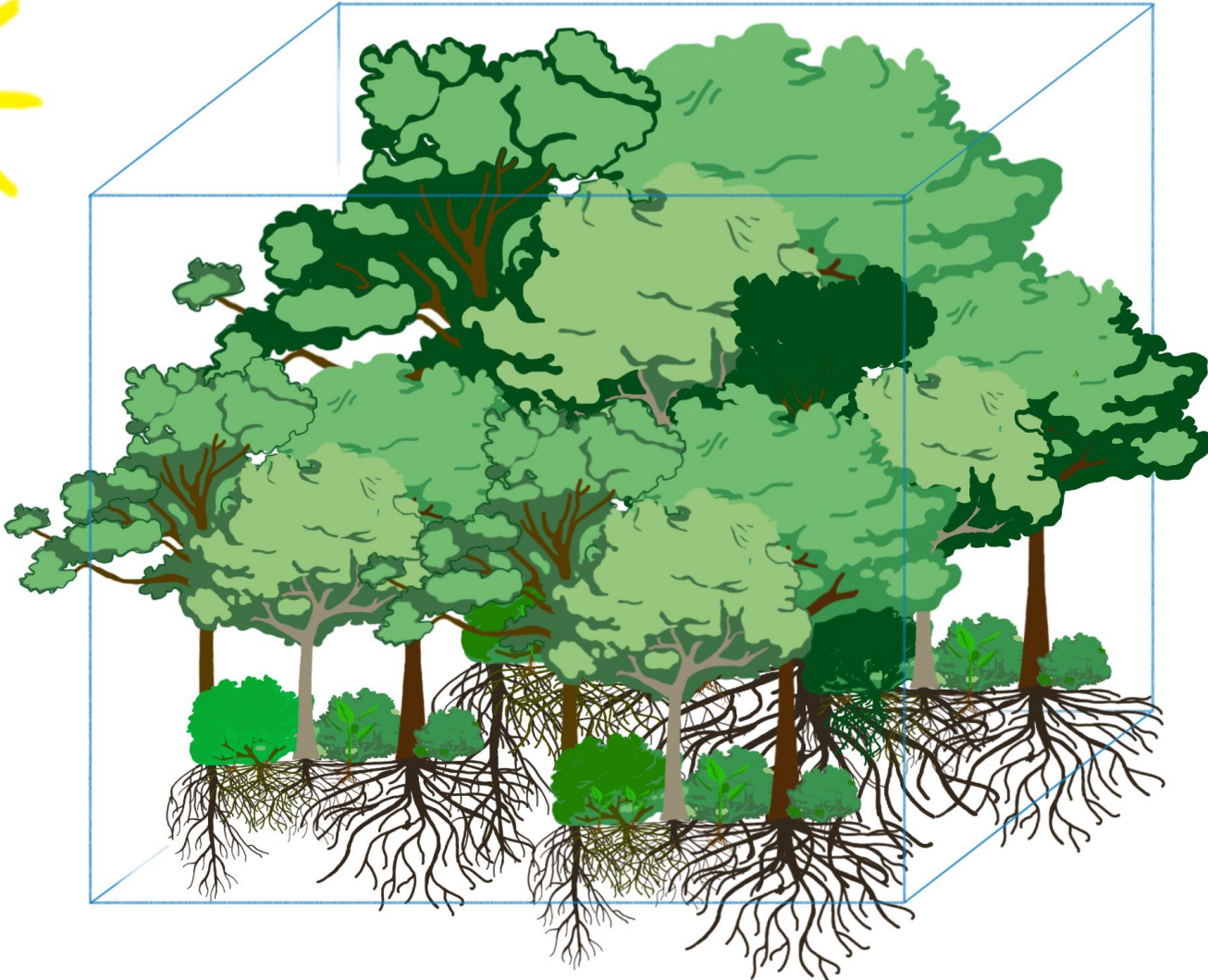


Introduction – Environmental space



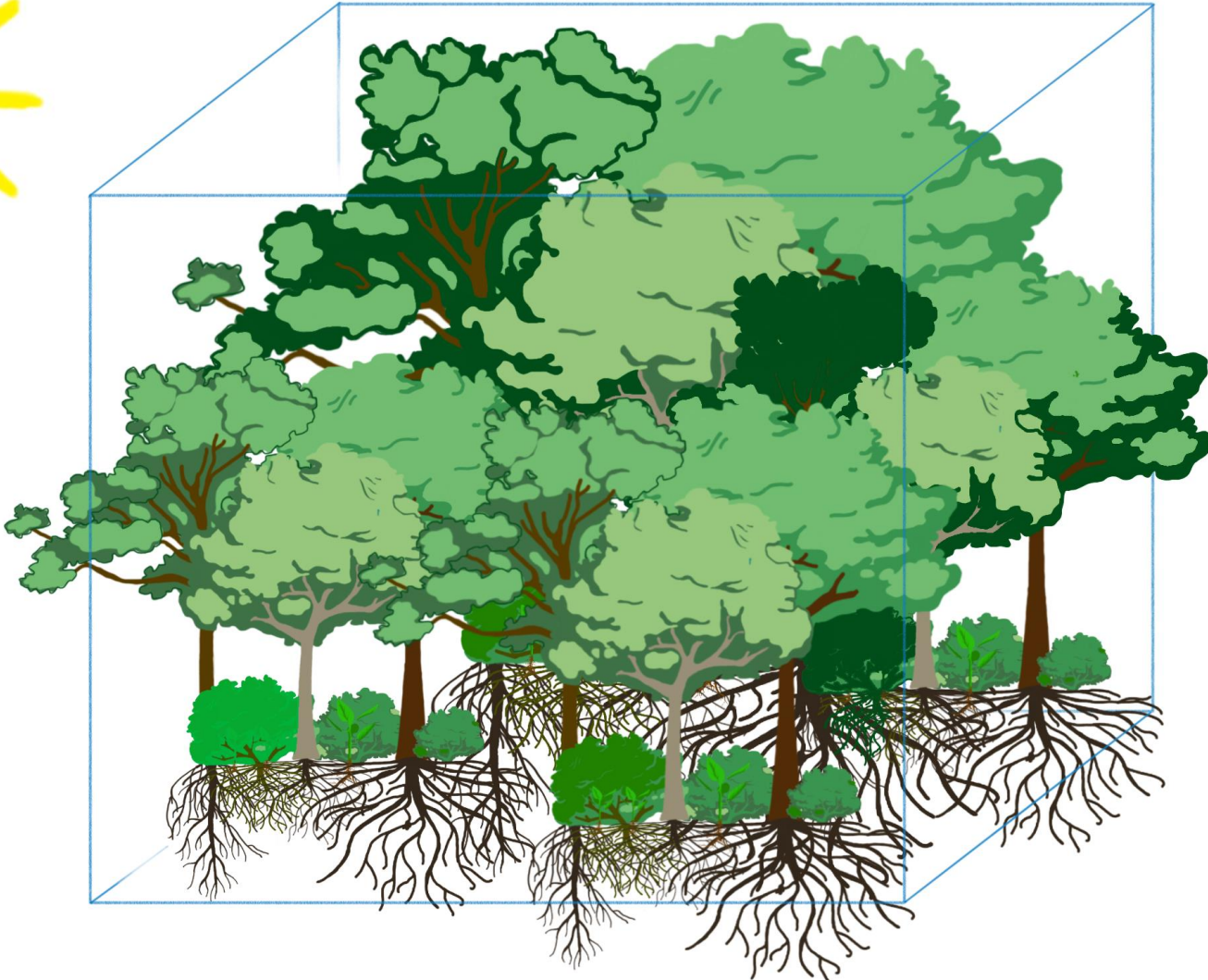
[1] Hutchinson 1978 [2] Gallego-Tévar et al. 2018

Introduction – Environmental space



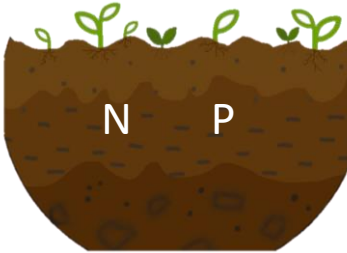
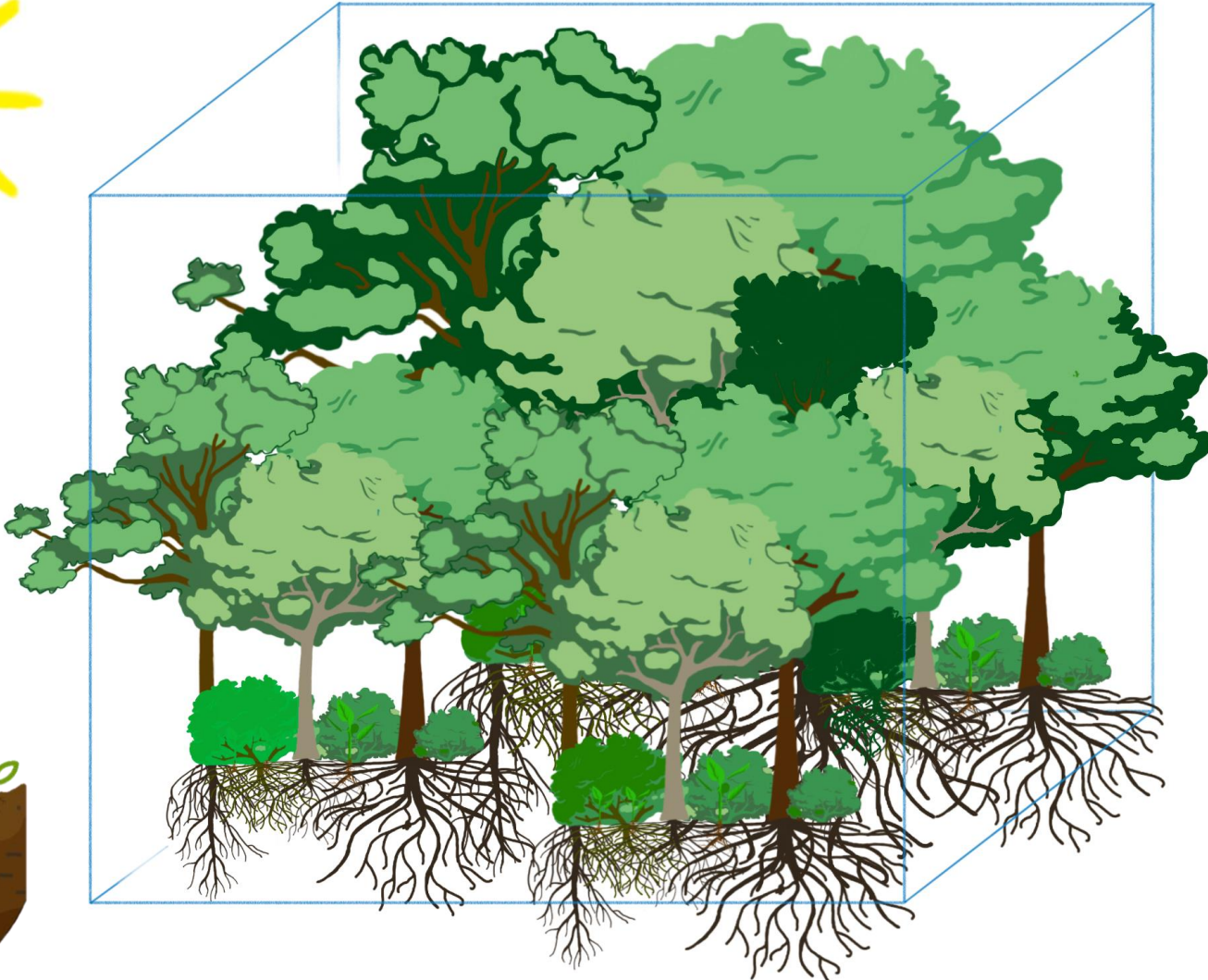
[1] Hutchinson 1978 [3] Reich et al. 2003

Introduction – Environmental space



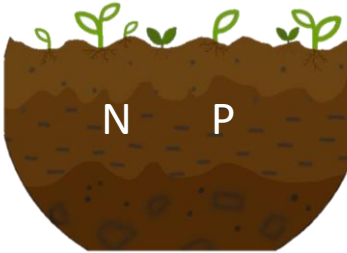
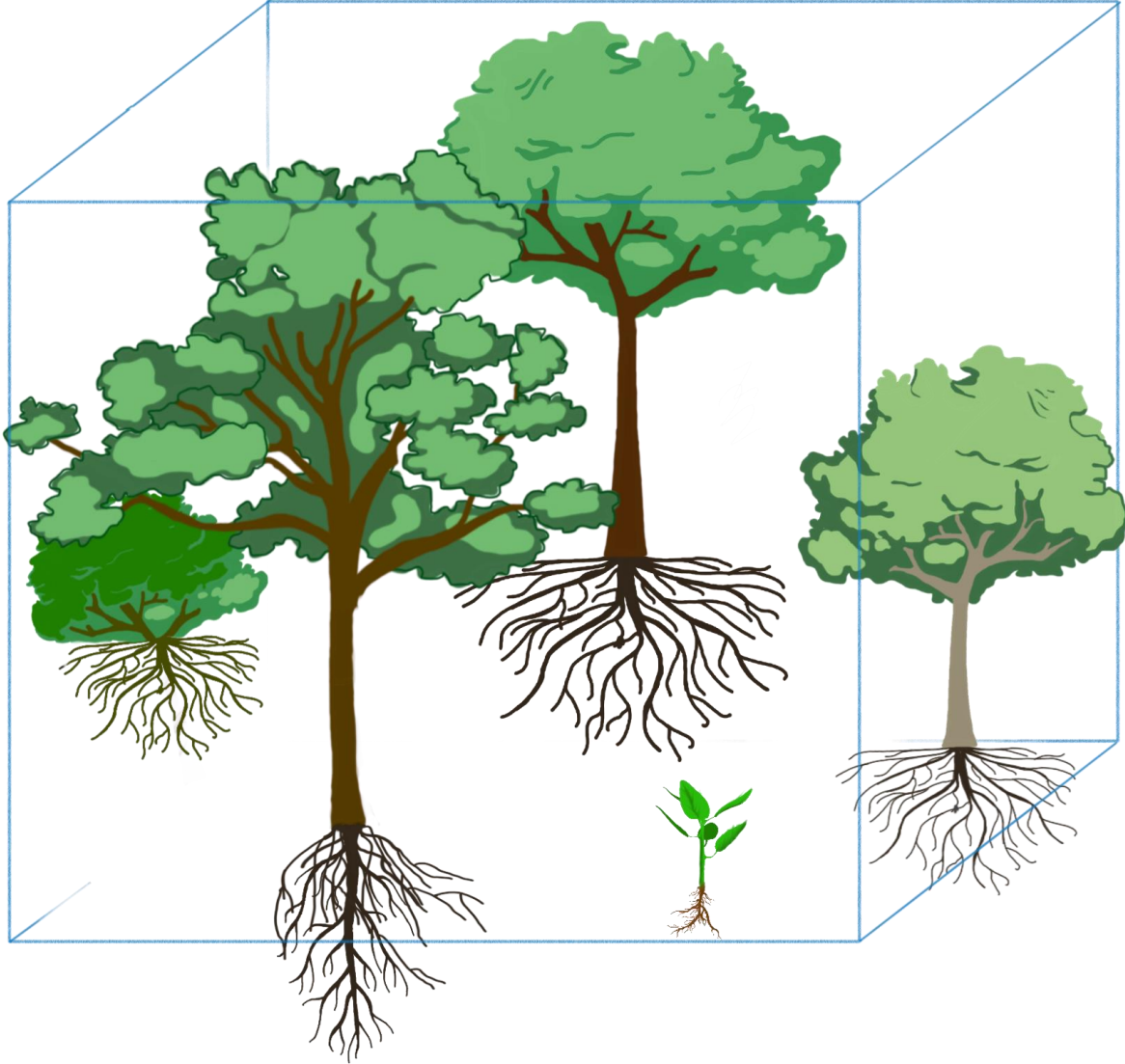
[1] Hutchinson 1978 [3] Reich et al. 2003

Introduction – Environmental space

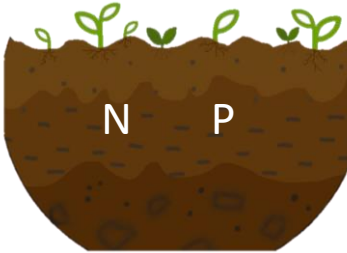
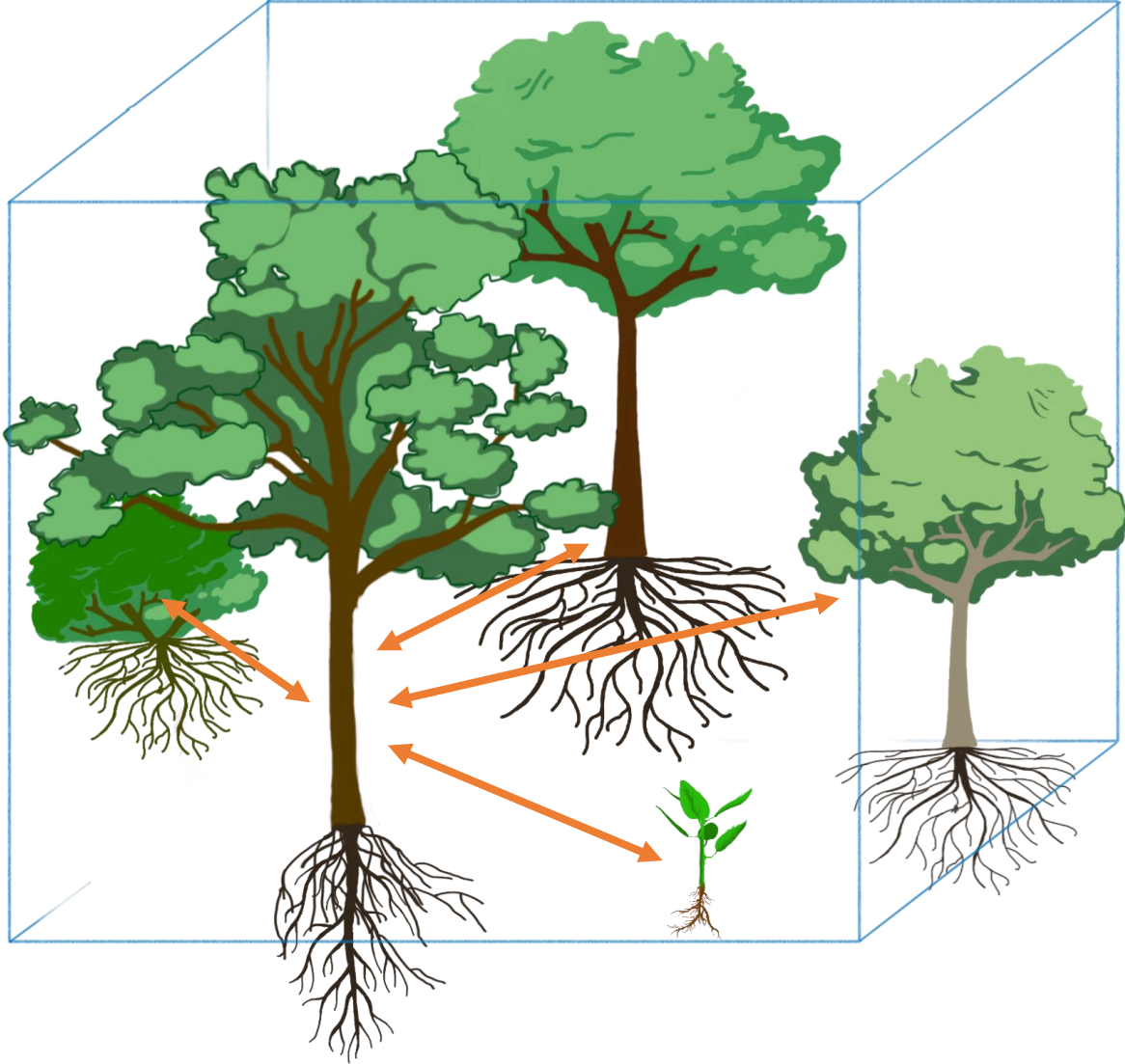


[1] Hutchinson 1978 [3] Reich et al. 2003 [4] García-Palacios et al. 2012

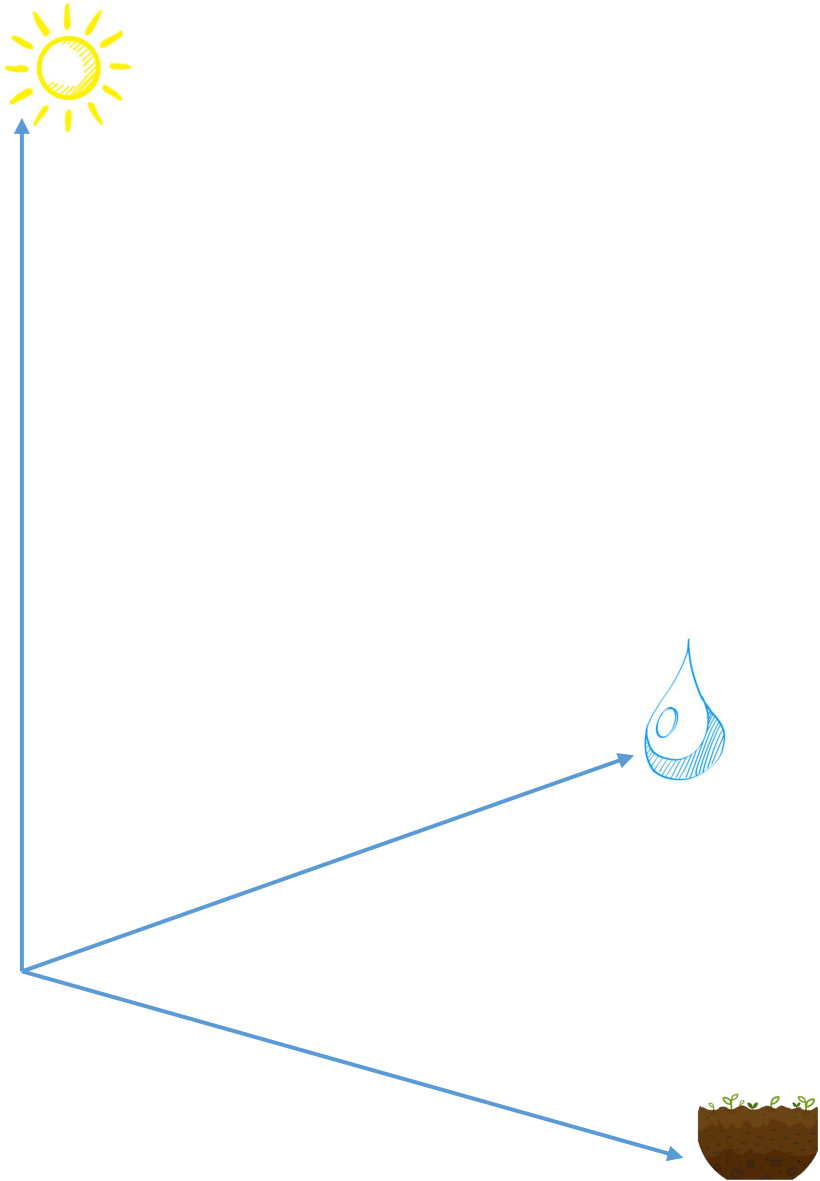
Introduction – Community assemblage



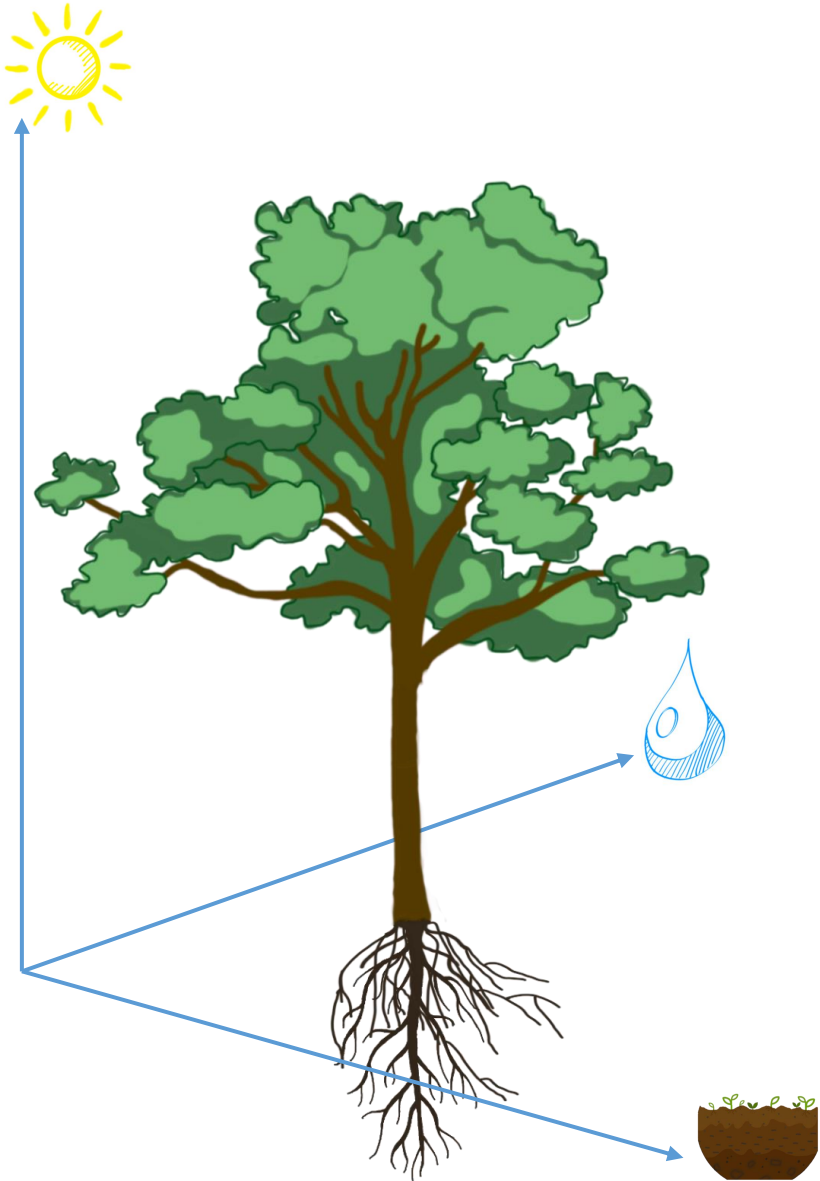
Introduction – Community assemblage



Introduction – Trait distances

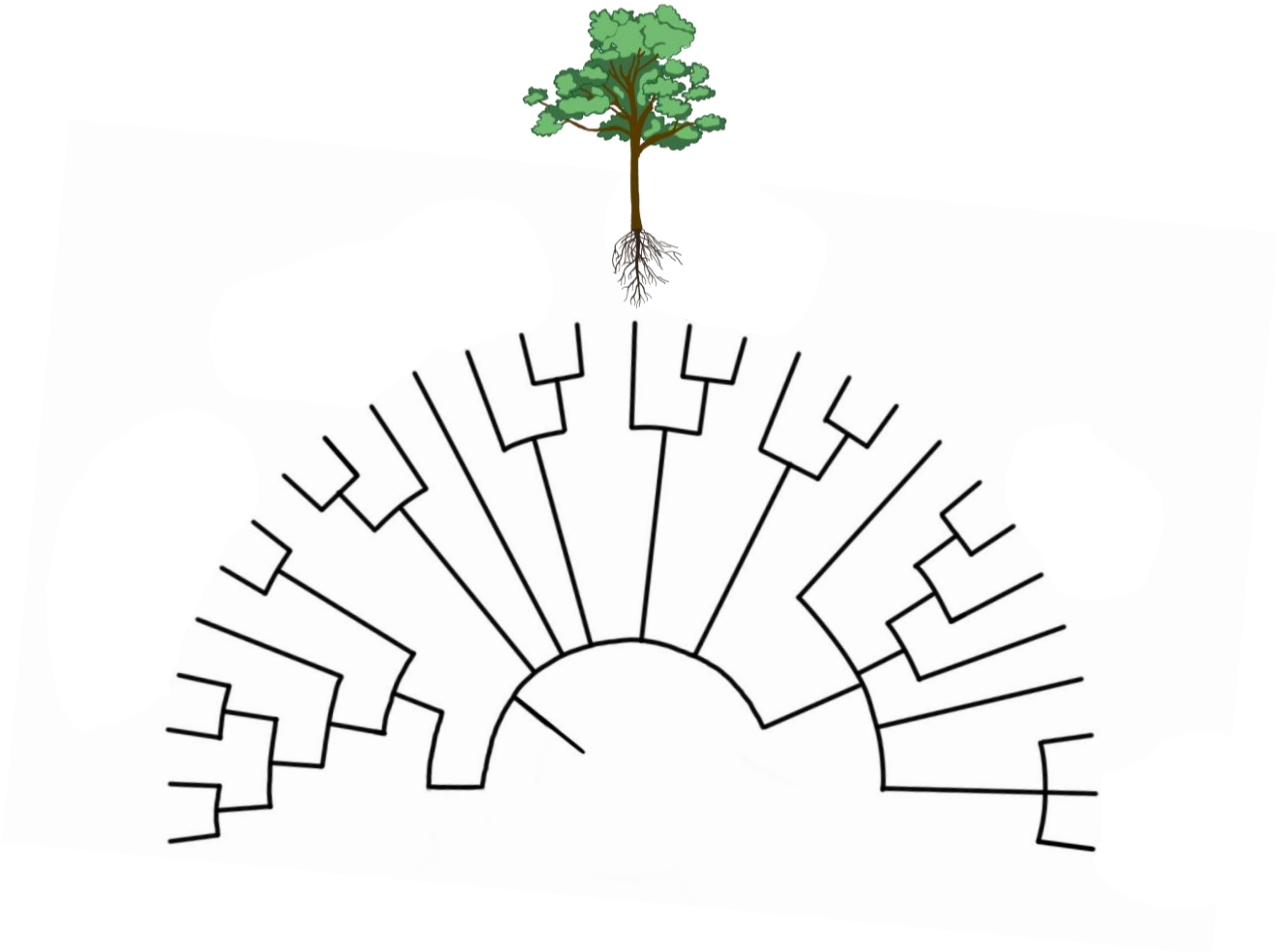
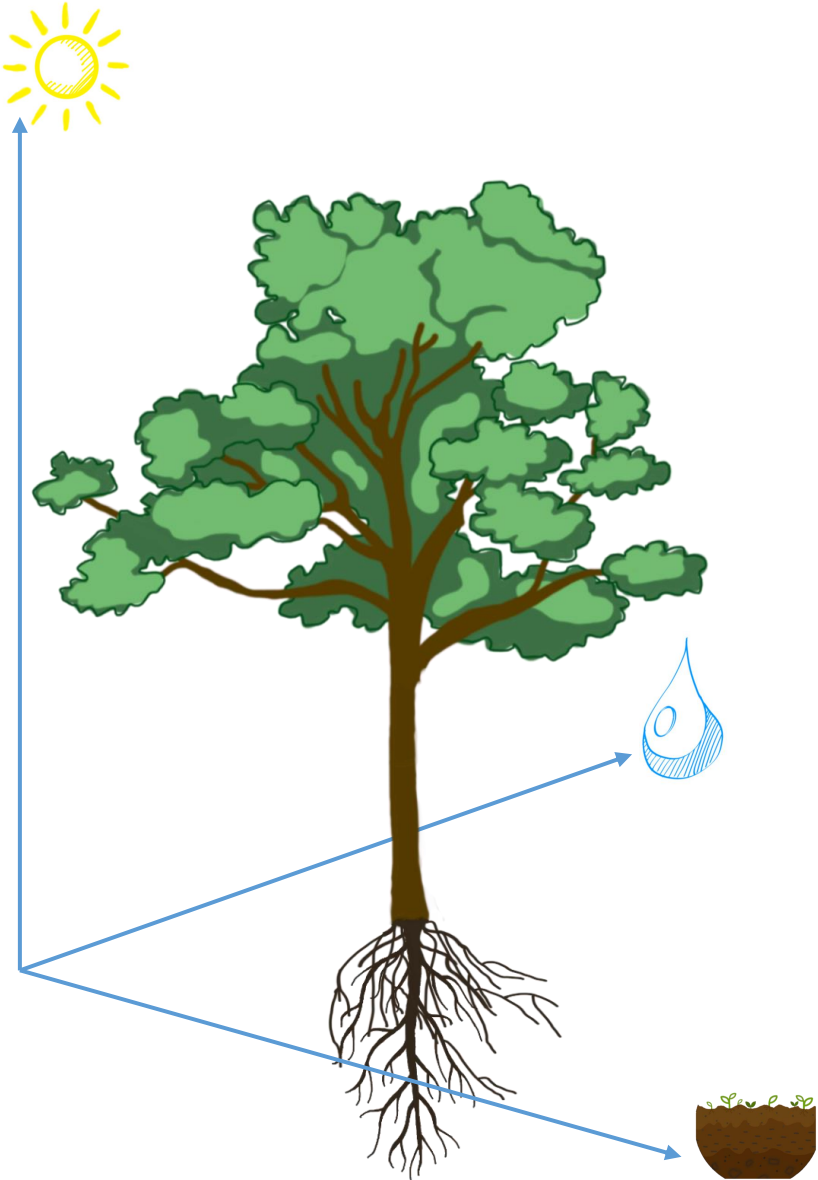


Introduction – Trait distances

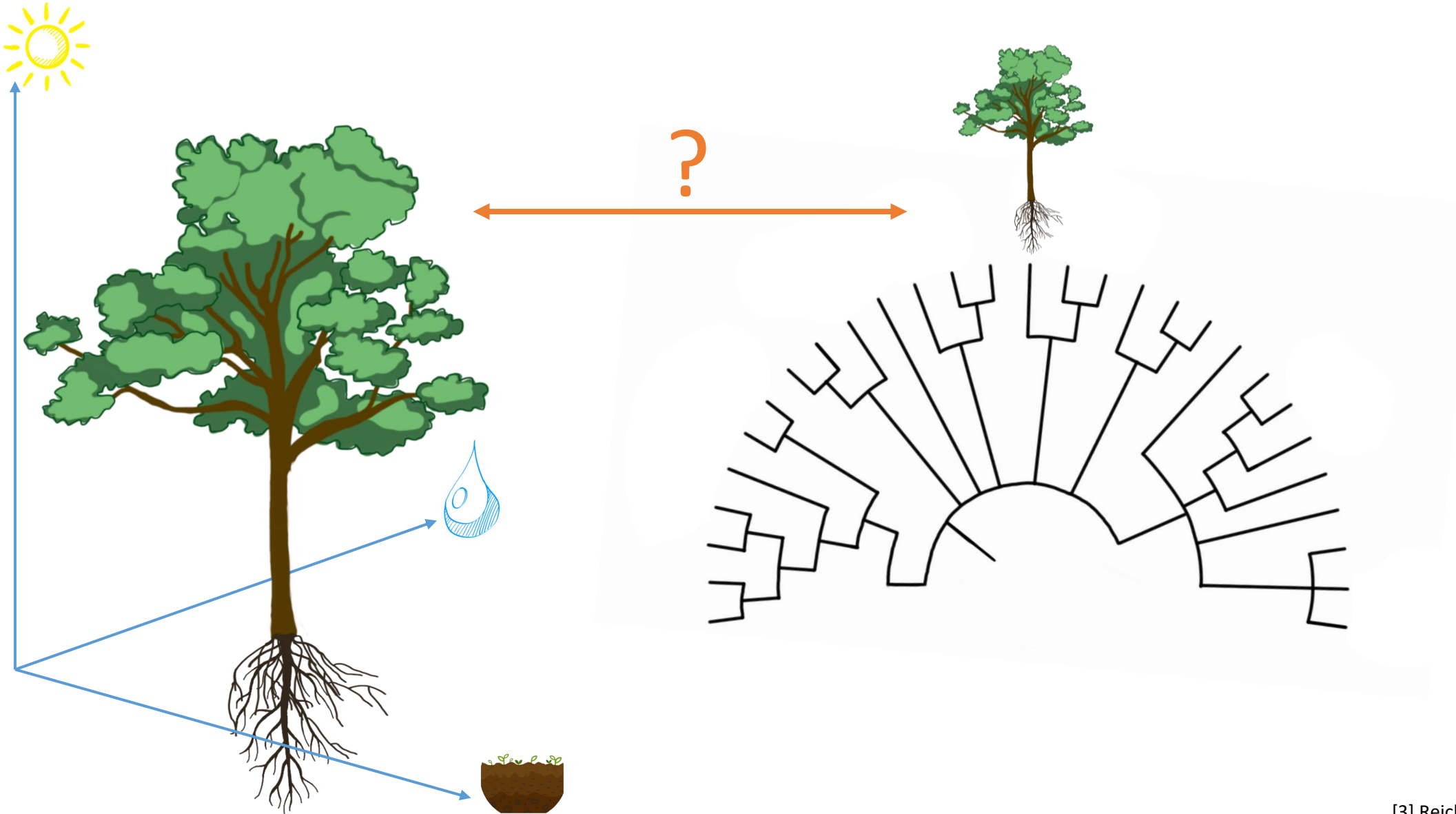


[5] McGill et al. 2006 [6] Wright et al. 2004

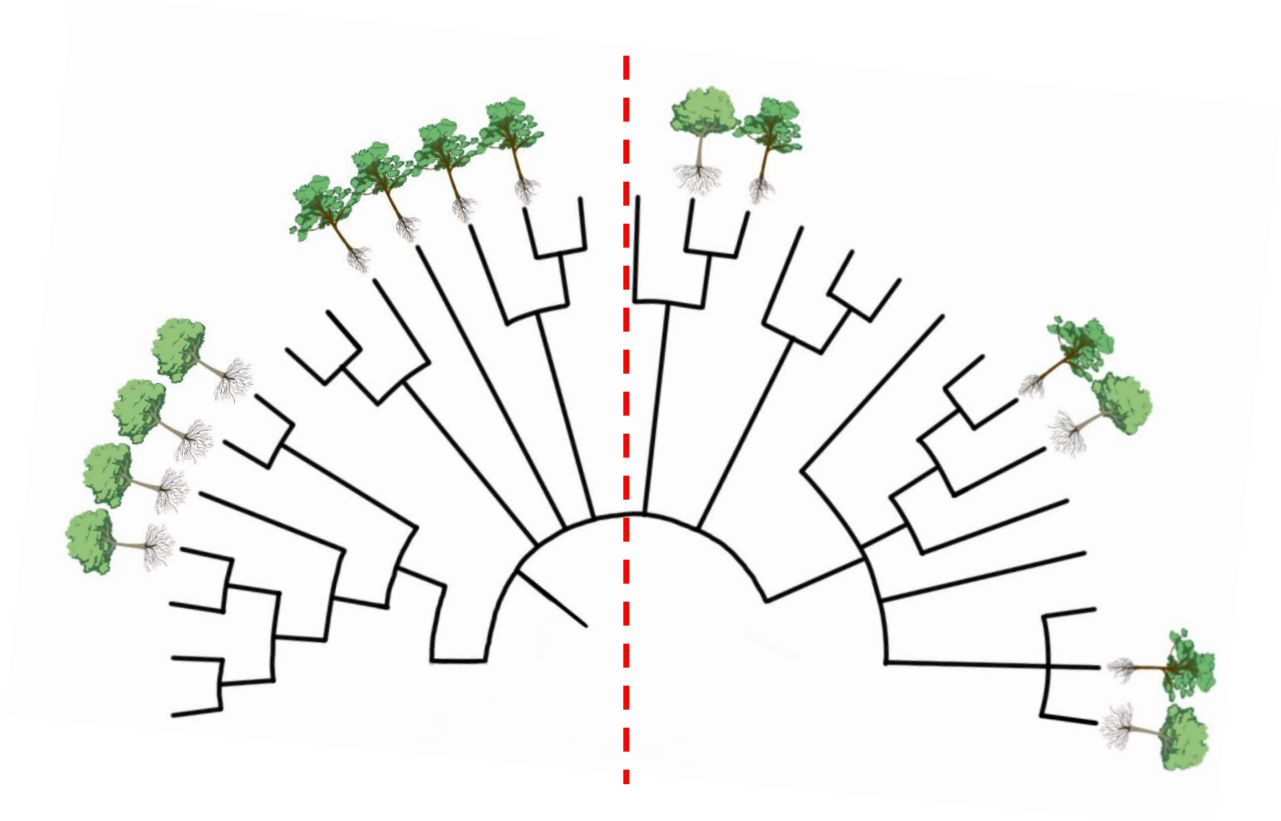
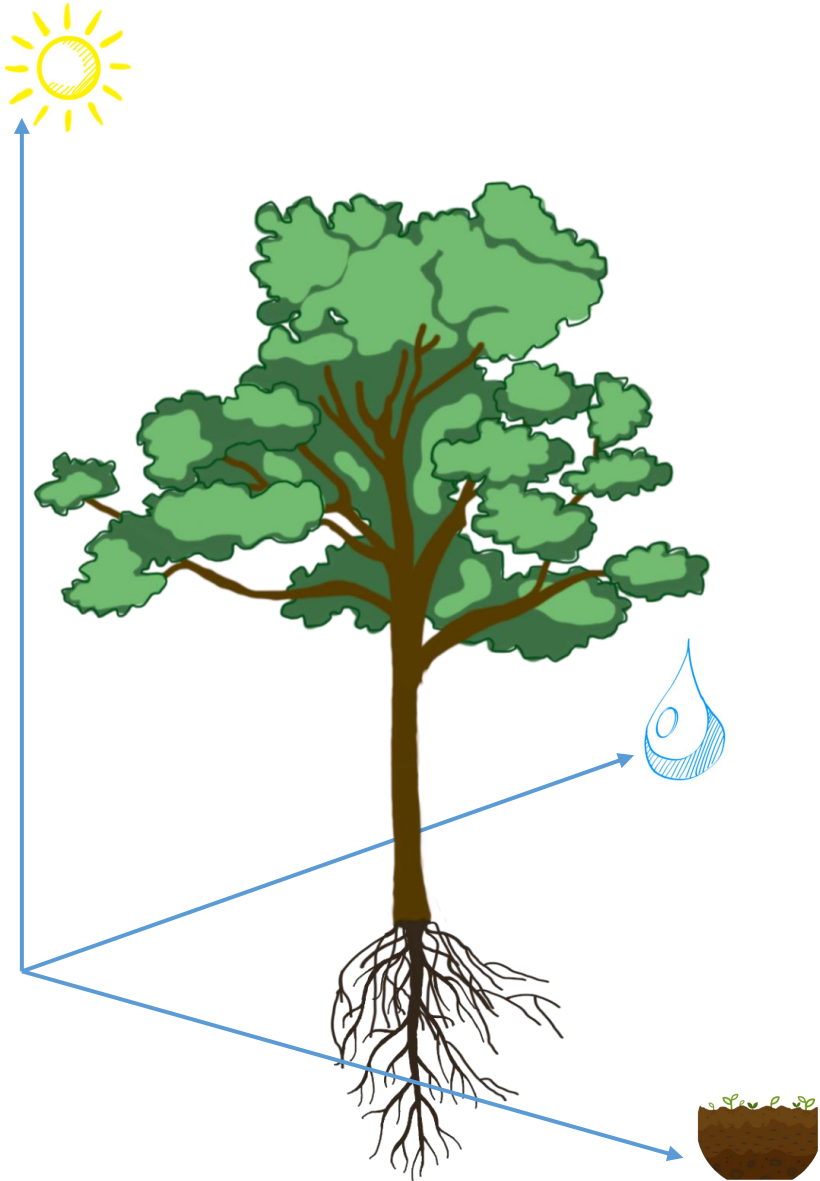
Introduction – Phylogenetic distances



Introduction – Phylogenetic distances

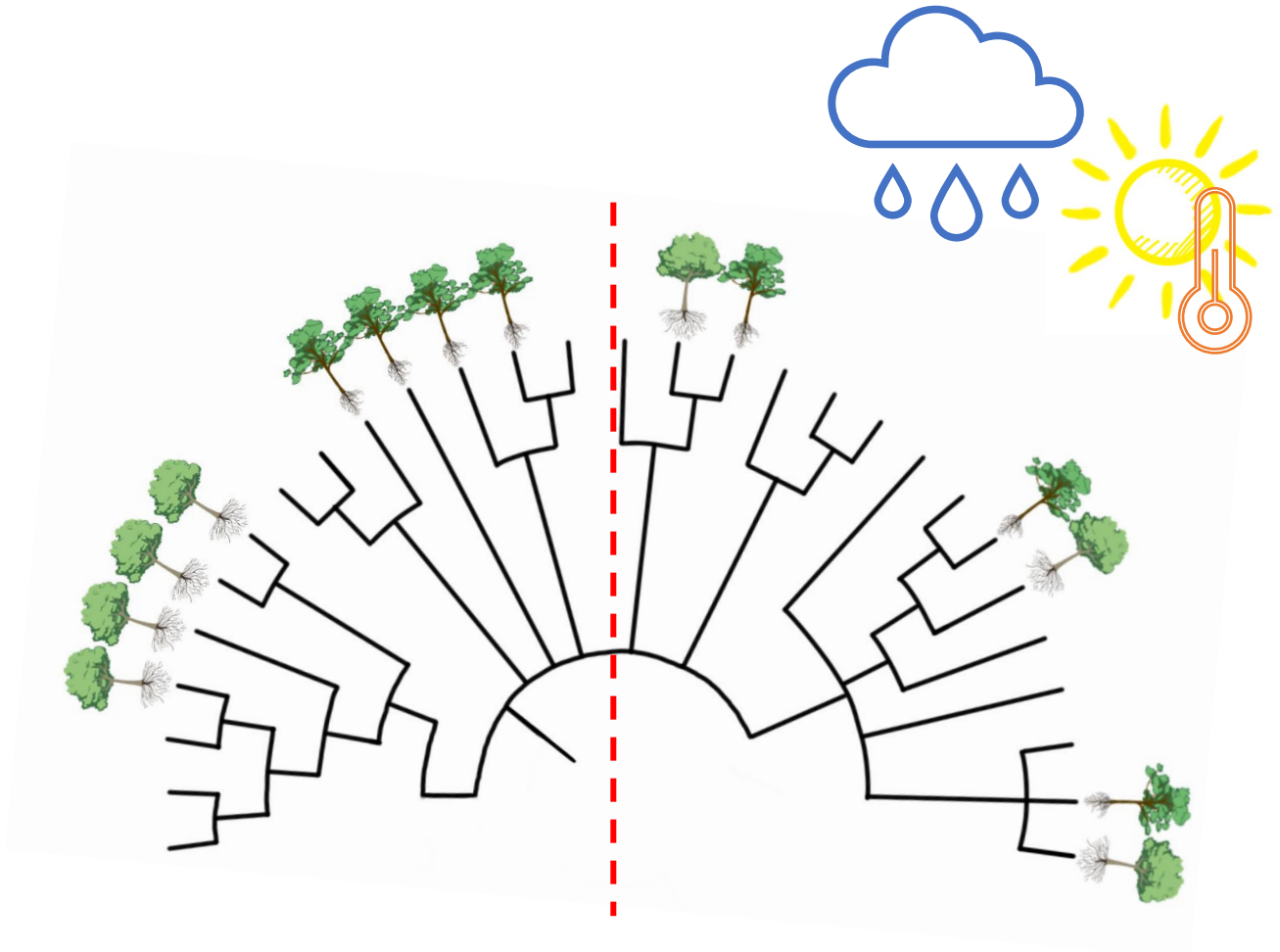
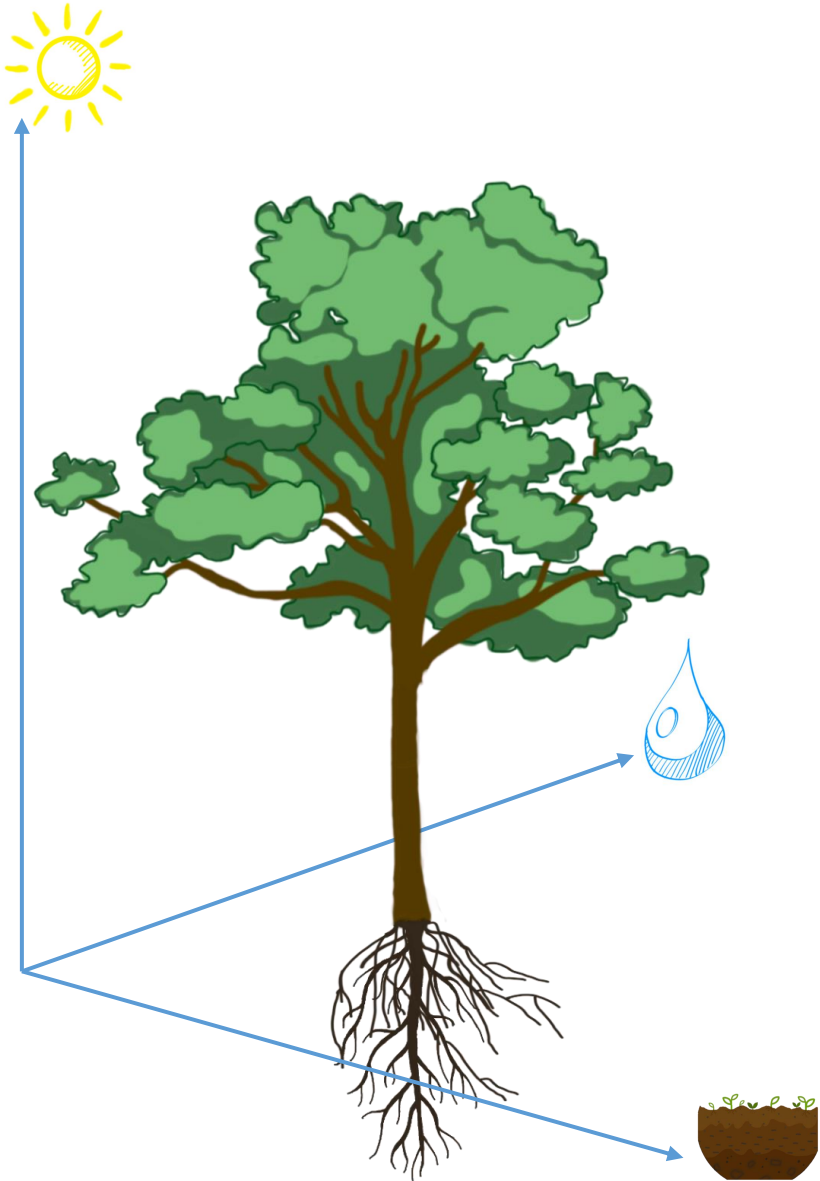


Introduction – Traits on the phylogeny



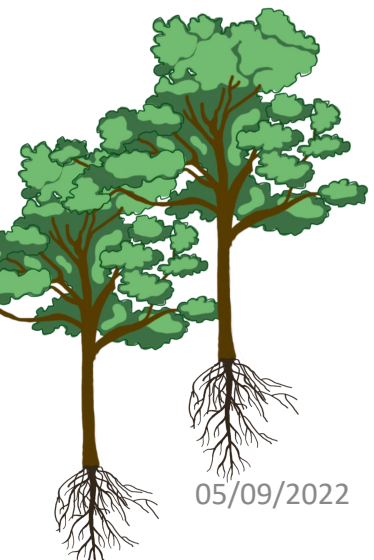
[7] Cavender-Bares et al. 2004 [8] Ackerly 2009 [9]

Introduction – Traits on the phylogeny



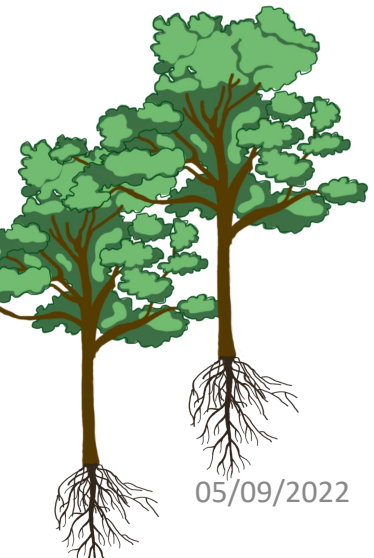
[9] Núñez-Farfán et al. 2007 [10] Castagneyrol et al. 2014

H1: Functional and phylogenetic diversity are related at the global scale.



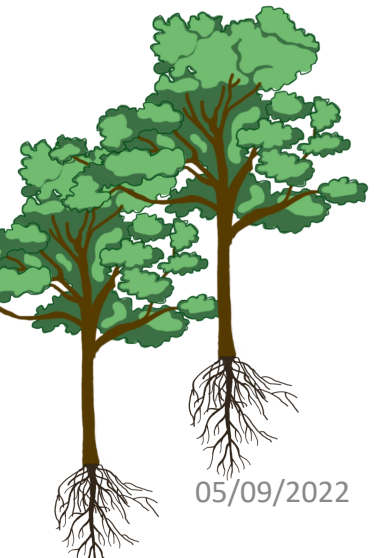
05/09/2022

- H1:** Functional and phylogenetic diversity are related at the global scale.
- H2:** Spatial patterns of functional and phylogenetic diversity differ from each other.

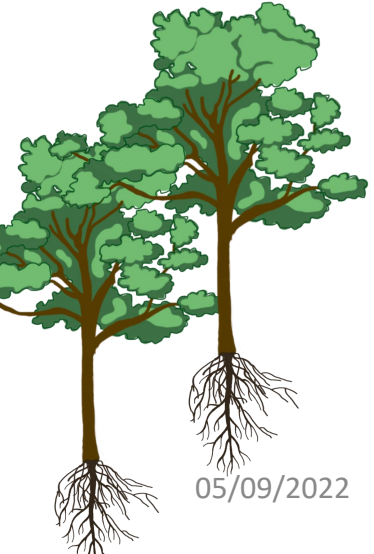


05/09/2022

- H1:** Functional and phylogenetic diversity are related at the global scale.
- H2:** Spatial patterns of functional and phylogenetic diversity differ from each other.
- H3:** Distribution pattern of functional diversity depends on current climatic conditions.



- H1:** Functional and phylogenetic diversity are related at the global scale.
- H2:** Spatial patterns of functional and phylogenetic diversity differ from each other.
- H3:** Distribution pattern of functional diversity depends on current climatic conditions.
- H4:** Spatial pattern of phylogenetic diversity depends on past climatic events, i.e. climatic conditions after the last glacial maximum.





sPlot

1,977,637 vegetation-plots



a project of  **iDiv**



sPlot

1,977,637 vegetation-plots^[11]

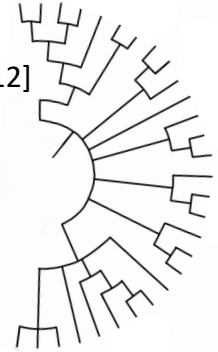
a project of



iDiv

GBOTB for seed plants^[12]

Clade in the phylogeny
for pteridophytes^[13]

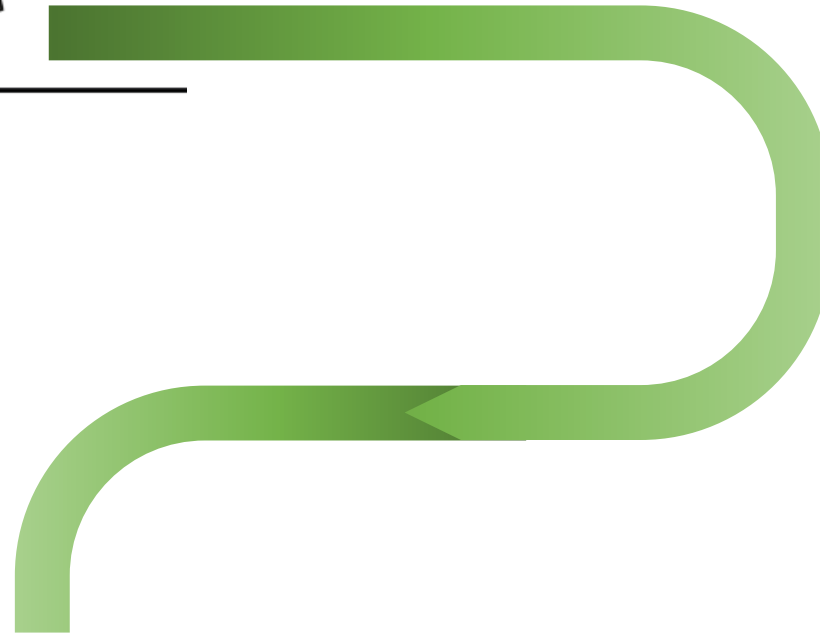


[11] Bruelheide et al. 2019 [12] Smith and Brown 2018 [13] Zanne et al. 2014



1,977,637 vegetation-plots^[11]

a project of  iDiv



GBOTB for seed plants^[12]

Clade in the phylogeny for pteridophytes^[13]




Specific leaf area, plant height and specific root length from the gap-filled TRY traits^[14, 15, 16, 17]

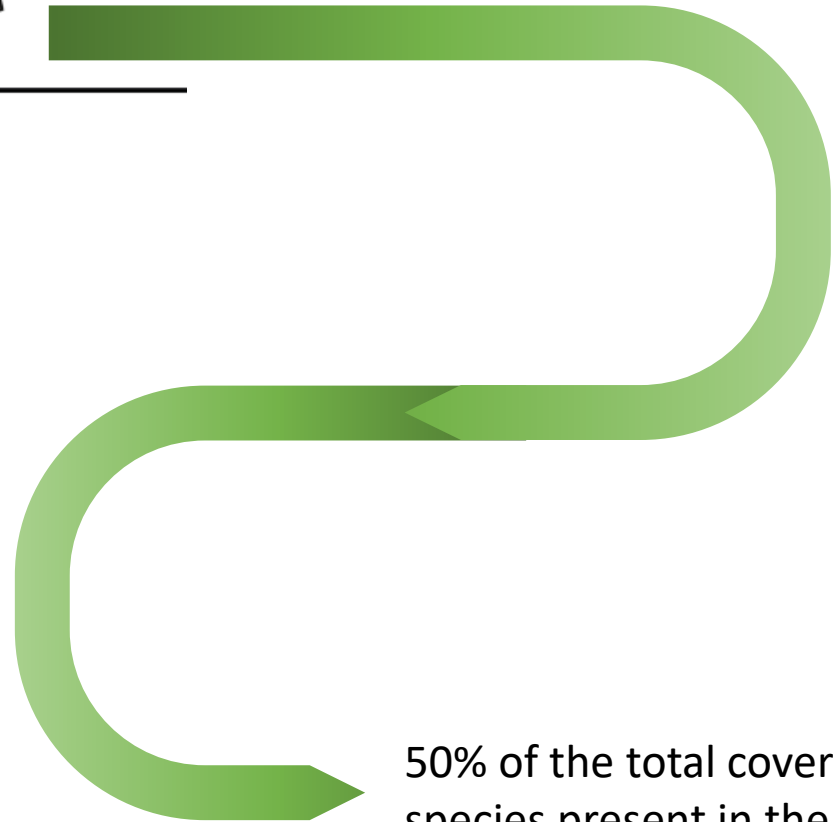
[11] Bruelheide et al. 2019 [12] Smith and Brown 2018 [13] Zanne et al. 2014 [14] Shan et al. 2012 [15] Fazayeli et al. 2014 [16] Schrodte et al. 2015 [17] Kattge et al. 2020



sPlot

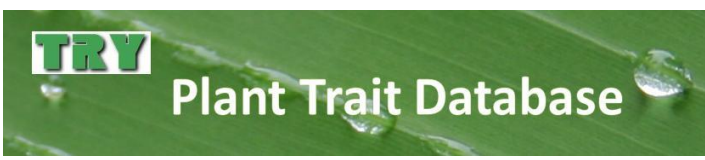
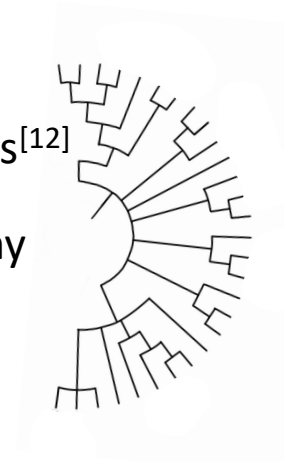
1,977,637 vegetation-plots^[11]

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Specific leaf area, plant height and
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50% of the total cover is represented by
species present in the phylogeny and traits

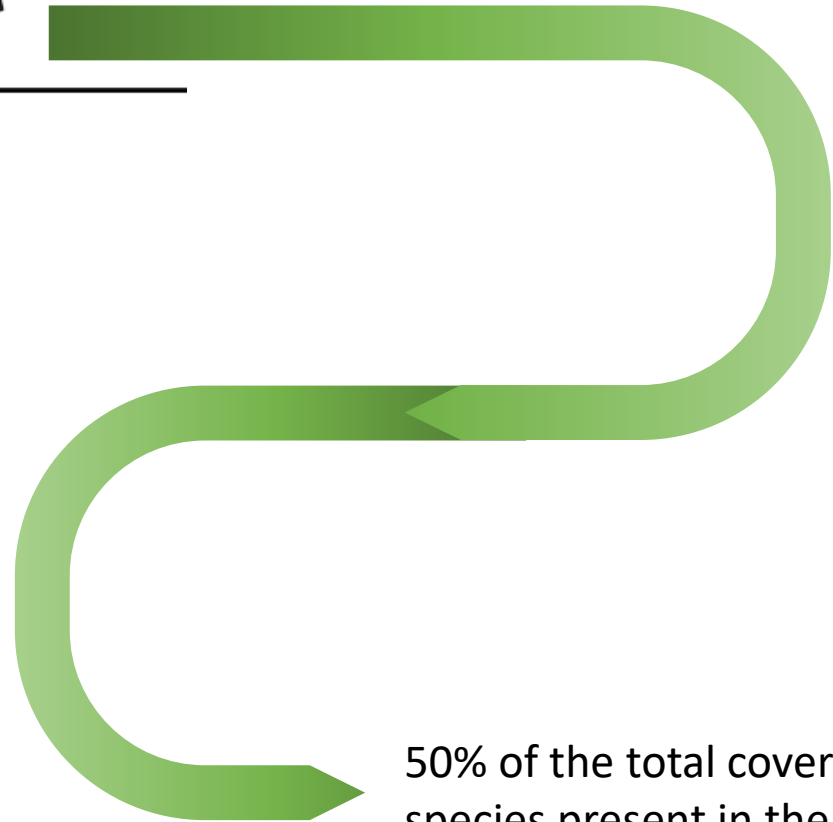
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Specific leaf area, plant height and
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→ 1,782,777 plots

[11] Bruelheide et al. 2019 [12] Smith and Brown 2018 [13] Zanne et al. 2014 [14] Shan et al. 2012 [15] Fazayeli et al. 2014 [16] Schrodte et al. 2015 [17] Kattge et al. 2020

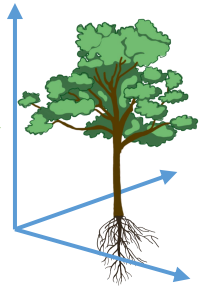
Methods – Response variables

1,782,777 plots



Methods – Response variables

1,782,777 plots



Functional diversity indices:

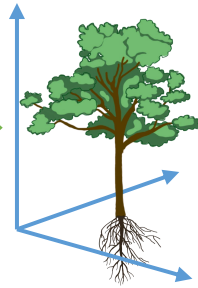
Rao's Quadratic Entropy (RQEF)^[18]

Functional dispersion (FDis)^[19]

[18] Rao 1982 [19] Anderson et al. 2006

Methods – Response variables

1,782,777 plots



Functional diversity indices:

Rao's Quadratic Entropy (RQEF)^[18]

Functional dispersion (FDis)^[19]



Phylogenetic diversity indices:

Rao's Quadratic Entropy (RQEP)^[18]

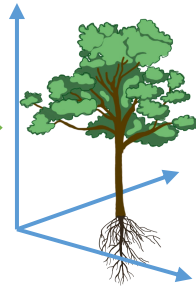
Mean pairwise distance (MPD)^[20]



[18] Rao 1982 [19] Anderson et al. 2006 [20] Webb et al. 2002

Methods – Response variables

1,782,777 plots



Functional diversity indices:

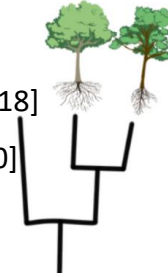
Rao's Quadratic Entropy (RQEF)^[18]

Functional dispersion (FDis)^[19]

Phylogenetic diversity indices:

Rao's Quadratic Entropy (RQEP)^[18]

Mean pairwise distance (MPD)^[20]



Correction for species richness dependencies^[21]

$$SES.MPD = \frac{MPD_{obs} - \overline{MPD}_{rand_{SR}}}{\sigma_{rand_{SR}}}$$



[18] Rao 1982 [19] Anderson et al. 2006 [20] Webb et al. 2002 [21] Botta-Dukát 2018



Recent climate conditions: 19 bioclimatic variables from CHELSA v.2.1^[22, 23]

Preselection with a principal component analyses

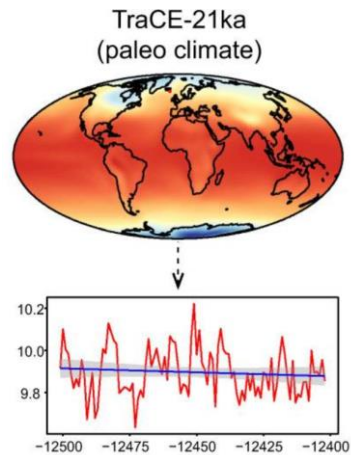
→ 5 climate variables



Recent climate conditions: 19 bioclimatic variables from CHELSA v.2.1^[22, 23]

Preselection with a principal component analyses

→ 5 climate variables

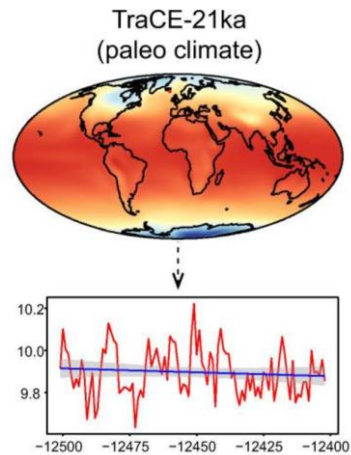


Worldwide stable climatic condition after the last glacial maximum (LGM) from StableClim v.1.1^[24]

[22] Karger et al. 2017 [23] Karger et al. 2018 [24] Brown et al. 2020



Recent climate conditions: 19 bioclimatic variables from CHELSA v.2.1^[22, 23]
Preselection with a principal component analyses
→ 5 climate variables



Worldwide stable climatic condition after the last glacial maximum (LGM) from StableClim v.1.1^[24]



a project of  iDiv

Plot size
Which plants were recorded
Categorization as forest or non-forest
Description of the vegetation type (biome)

[22] Karger et al. 2017 [23] Karger et al. 2018 [24] Brown et al. 2020

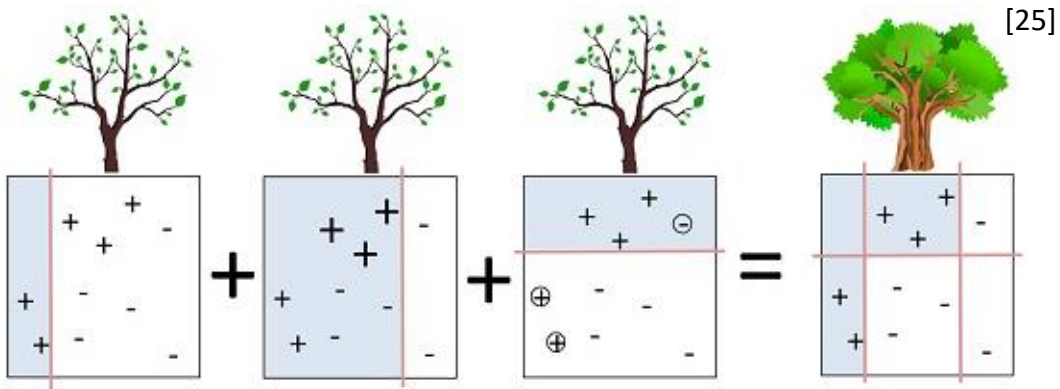
```
gam( SES.RQEF ~ SES.RQEP + s(Longitude, Latitude, bs = "sos"), family = "gaussian", method = "REML")
```

```
gam( SES.RQEF ~ SES.RQEP + s(Longitude, Latitude, bs = "sos"), family = "gaussian", method = "REML")
```

Boosted regression trees (BRT) to select explanatory variables with most relevant influence

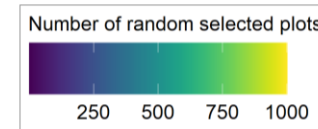
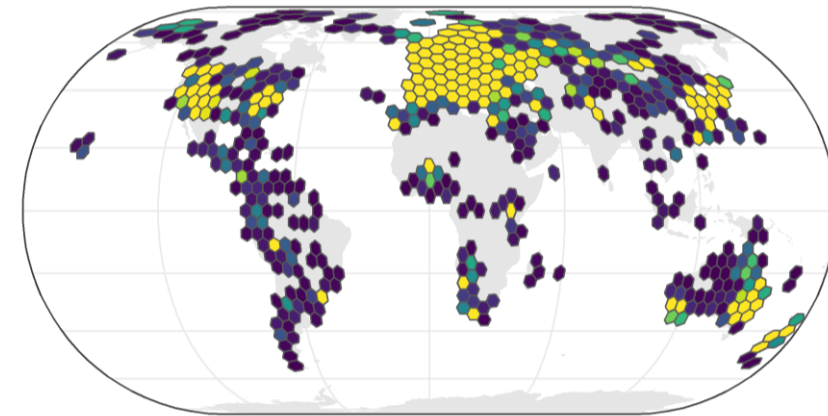
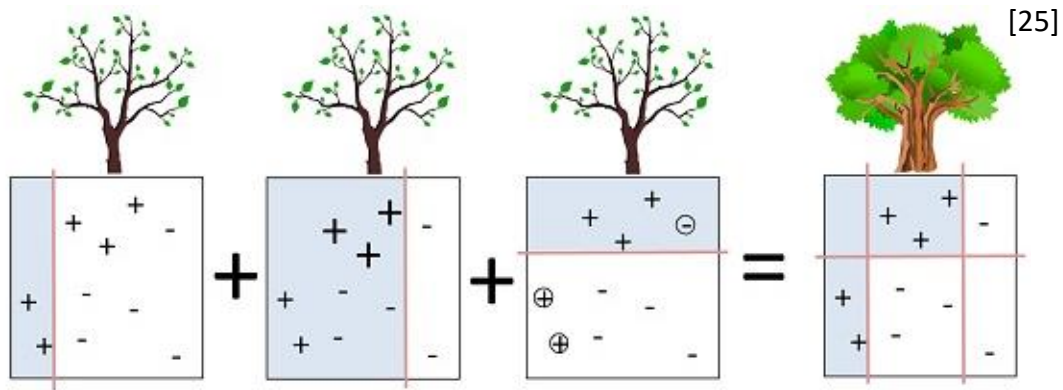
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gam( SES.RQEF ~ SES.RQEP + s(Longitude, Latitude, bs = "sos"), family = "gaussian", method = "REML")
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Boosted regression trees (BRT) to select explanatory variables with most relevant influence



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```

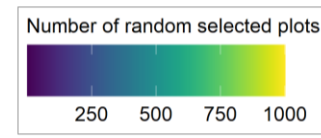
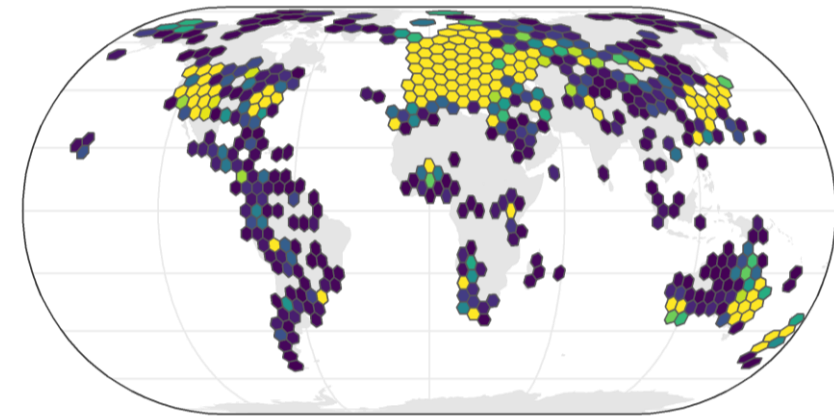
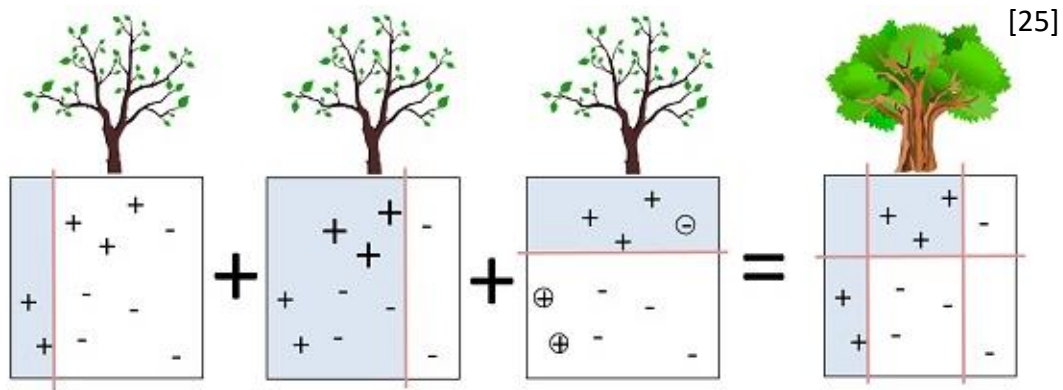
Boosted regression trees (BRT) to select explanatory variables with most relevant influence



Grid-cell size of ~209,000 km²

```
gam( SES.RQEF ~ SES.RQEP + s(Longitude, Latitude, bs = "sos"), family = "gaussian", method = "REML")
```

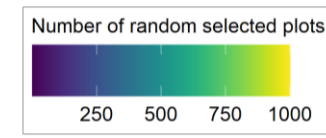
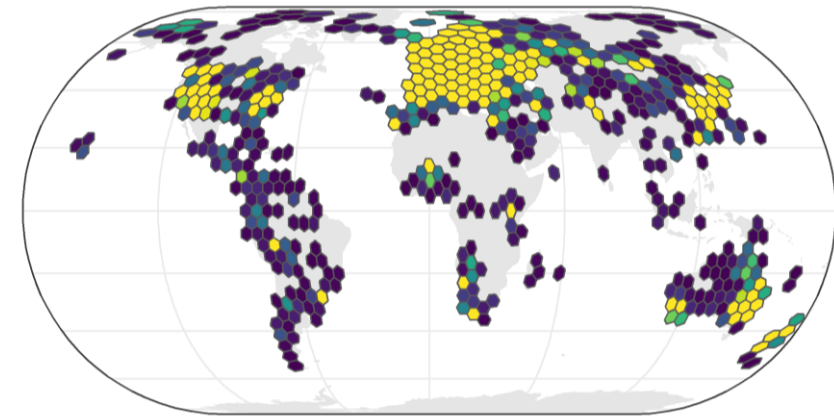
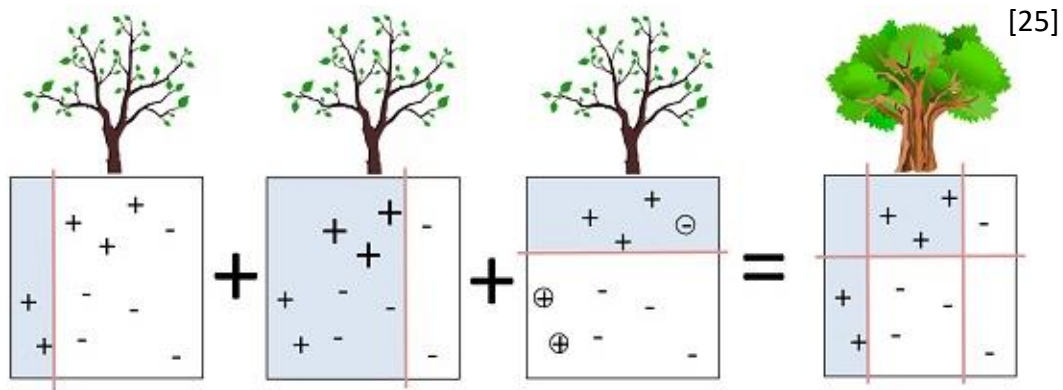
Boosted regression trees (BRT) to select explanatory variables with most relevant influence



Grid-cell size of $\sim 209,000 \text{ km}^2$
→ 181,151 plots per run

`gam(SES.RQEF ~ SES.RQEP + s(Longitude, Latitude, bs = "sos"), family = "gaussian", method = "REML")`

Boosted regression trees (BRT) to select explanatory variables with most relevant influence

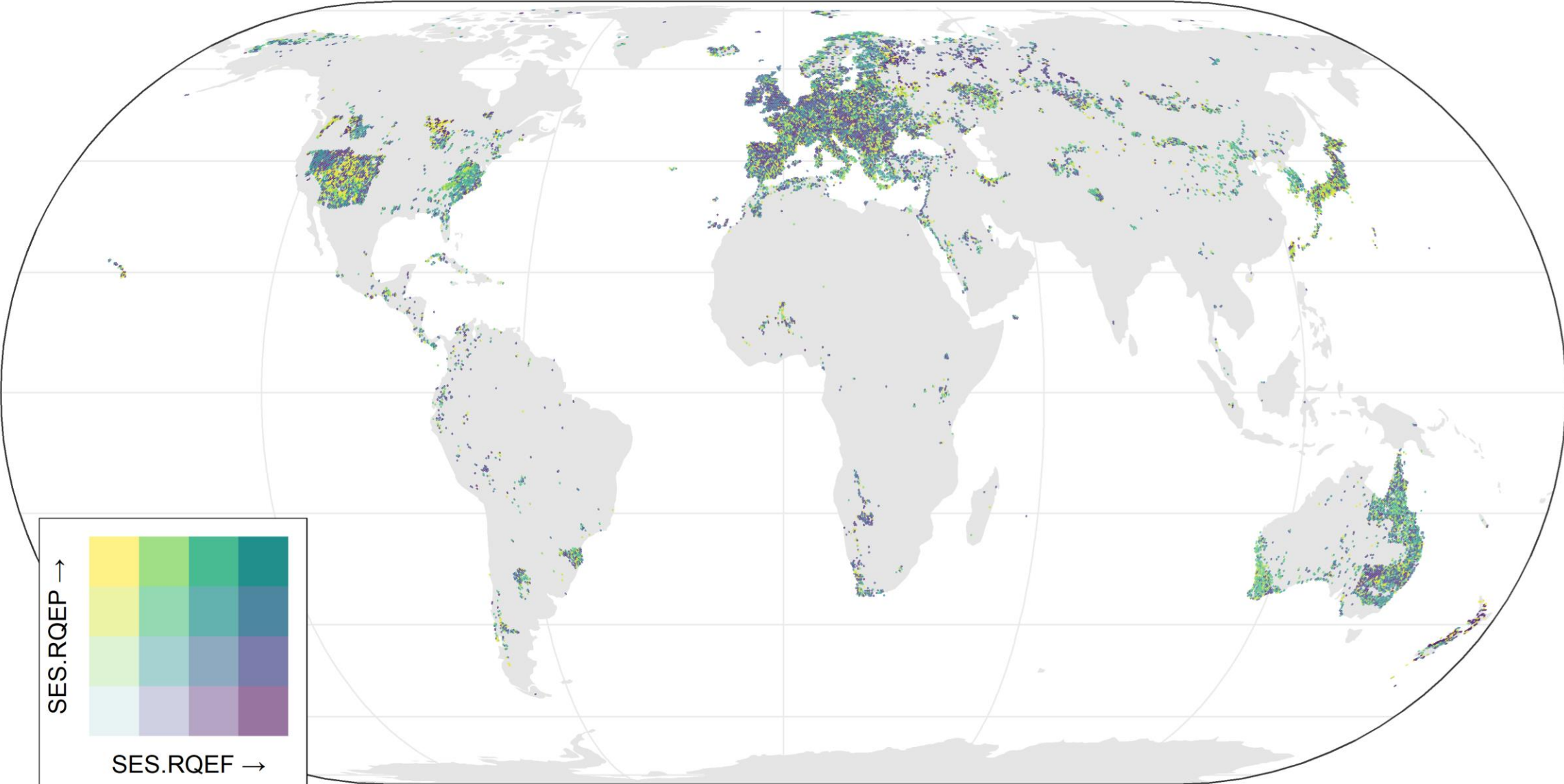


Grid-cell size of $\sim 209,000 \text{ km}^2$
 $\rightarrow 181,151$ plots per run

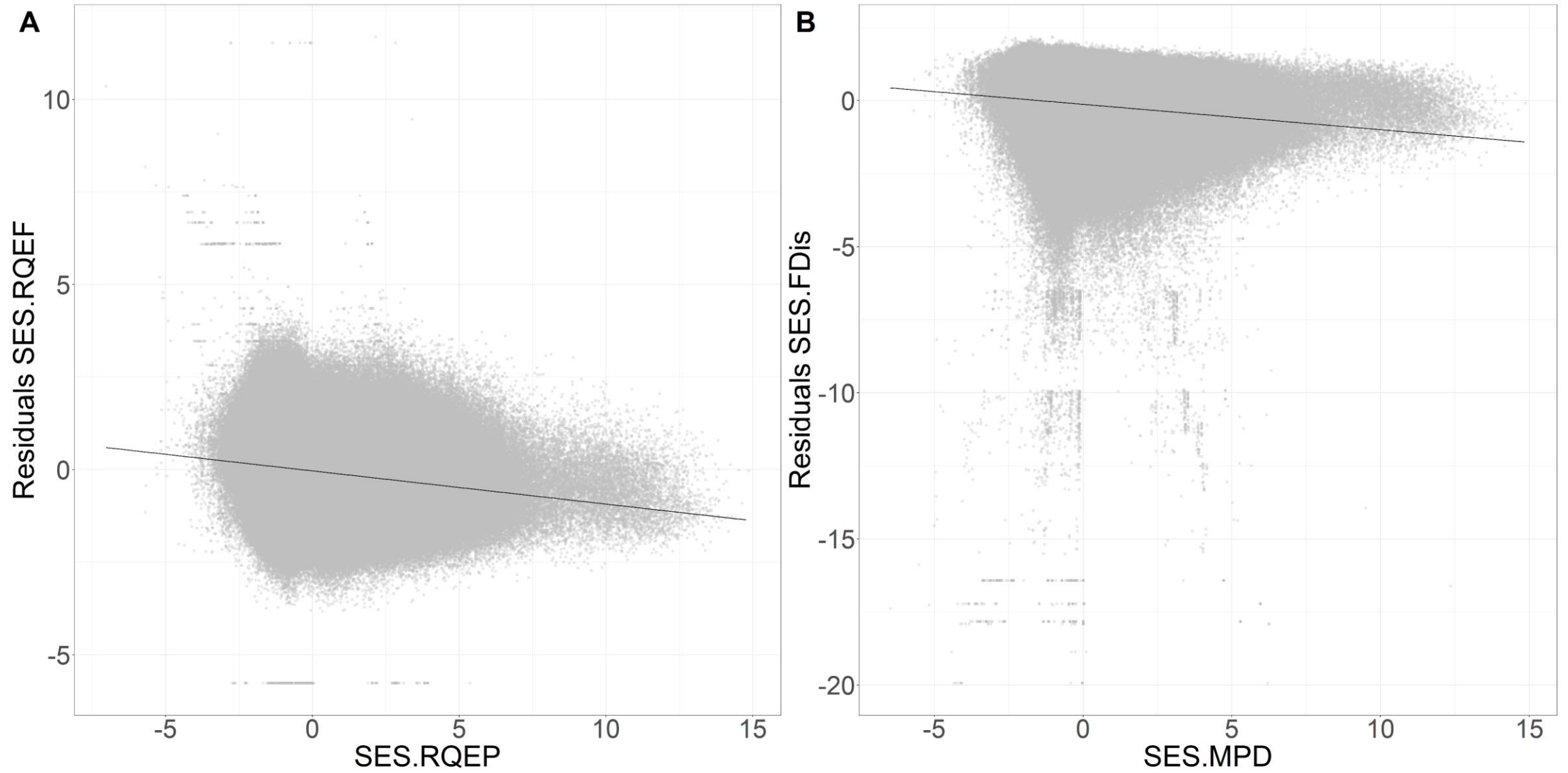
`gam(SES.RQEF ~ explanatory variables + s(Longitude, Latitude, bs = "sos"), family = "gaussian", method = "REML")`

[25] Colin et al. 2018

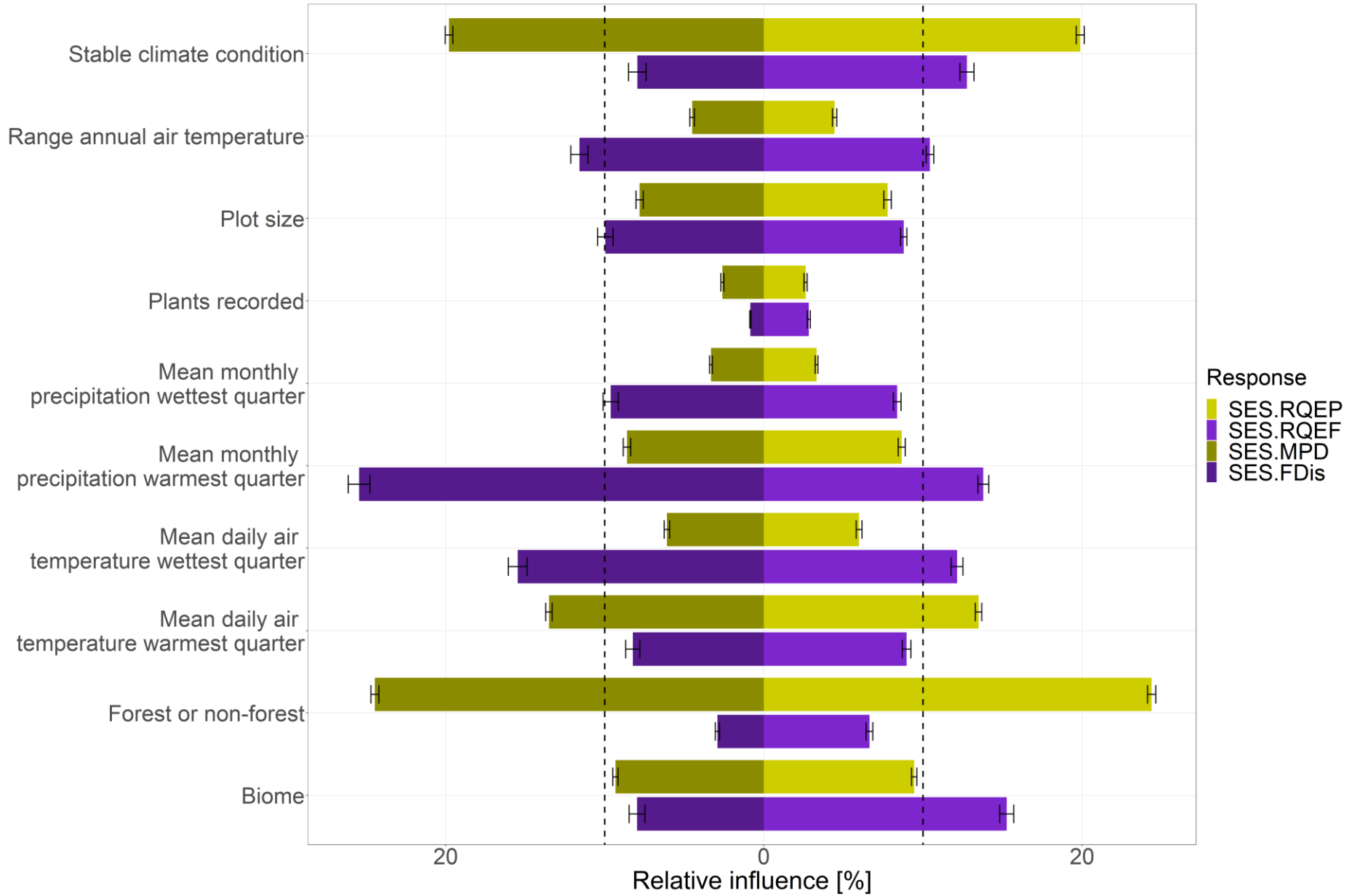
Results – Functional and phylogenetic diversity



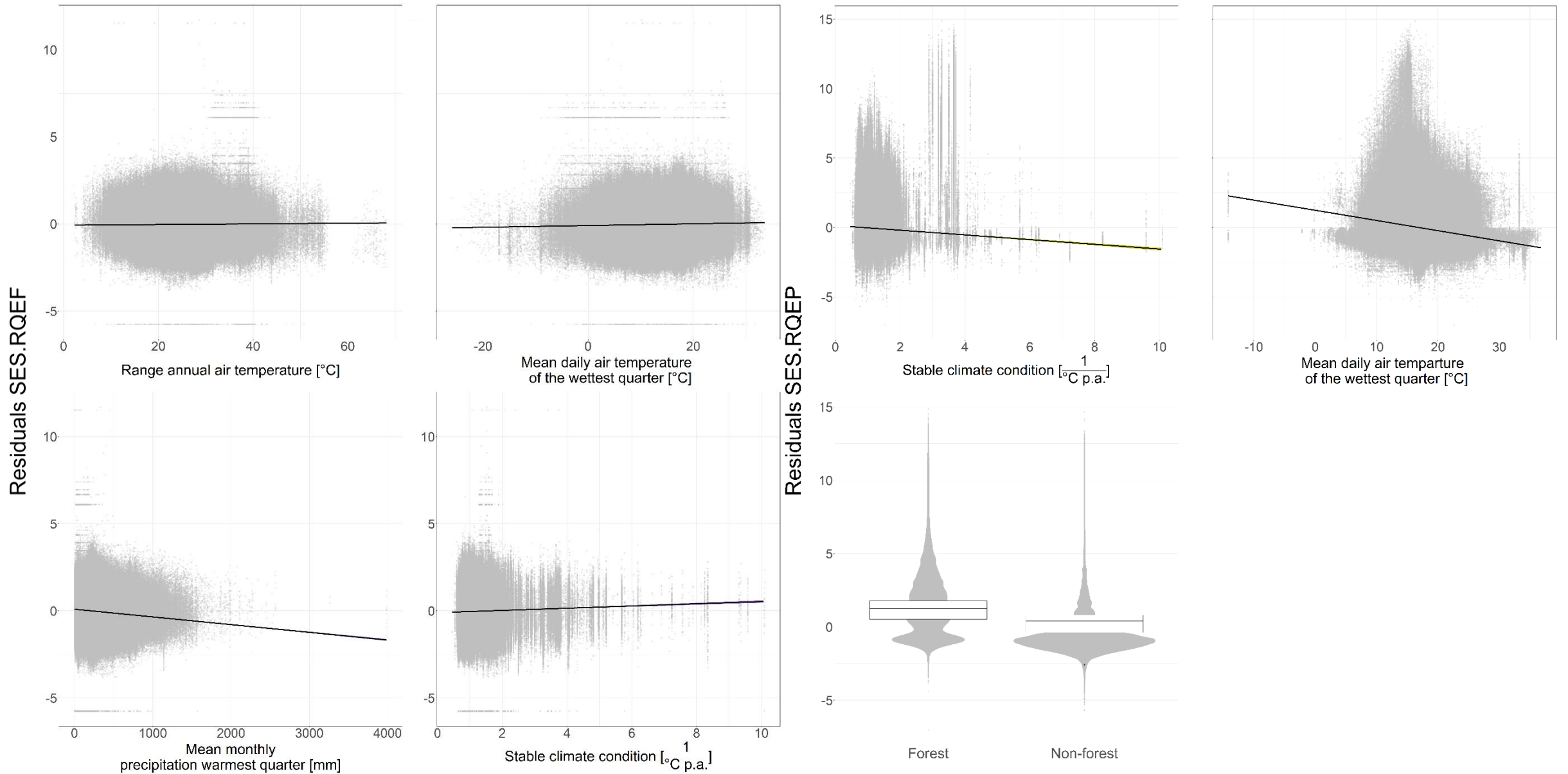
Results – Functional and phylogenetic diversity



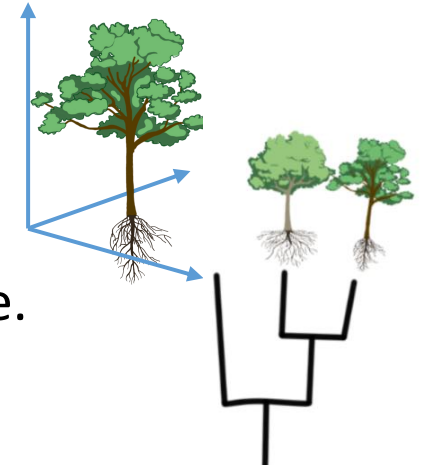
Results – Drivers of plant diversity



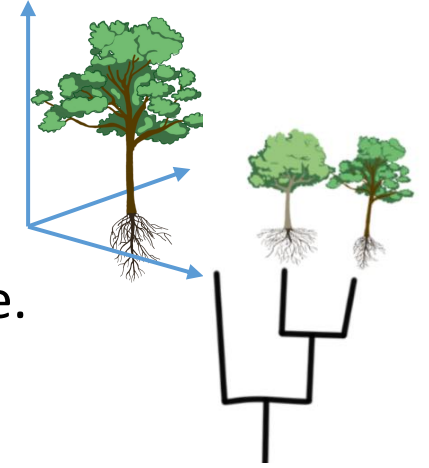
Results – Drivers of plant diversity

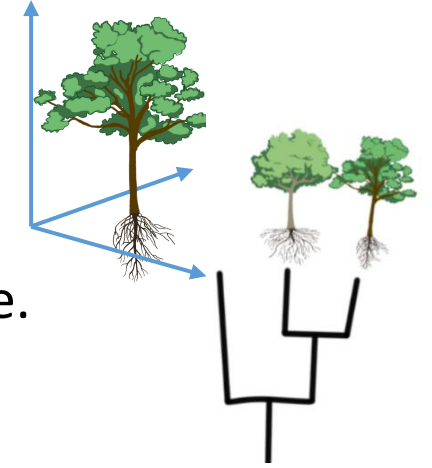


H1: Functional and phylogenetic diversity are related at the global scale.



H1: Functional and phylogenetic diversity are related at the global scale.
→ negatively correlated / tendency of trait overdispersion



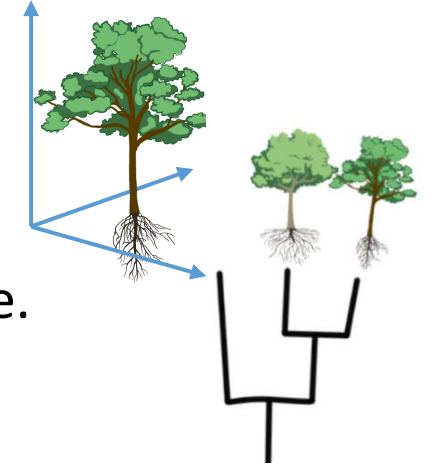


H1: Functional and phylogenetic diversity are related at the global scale.
→ negatively correlated / tendency of trait overdispersion

Positive correlation was shown before^[26]

Increasing with higher number of traits

[26] Tucker et al. 2018 [7] Cavender-Bares et al. 2004 [8] Ackerly 2009



H1: Functional and phylogenetic diversity are related at the global scale.
→ negatively correlated / tendency of trait overdispersion

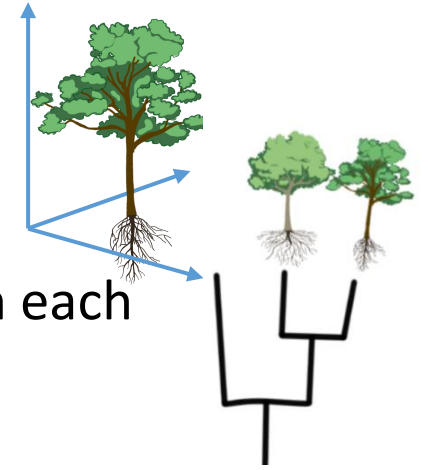
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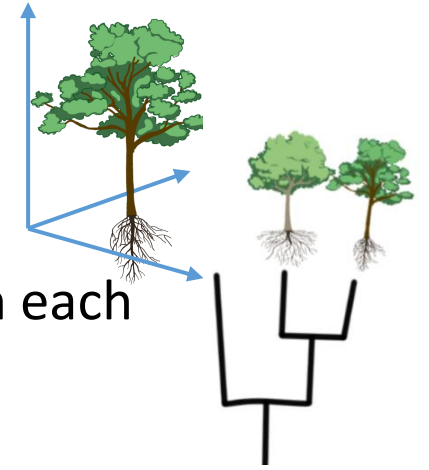
Traits map differently on the phylogeny at smaller spatial extent^[7, 8]

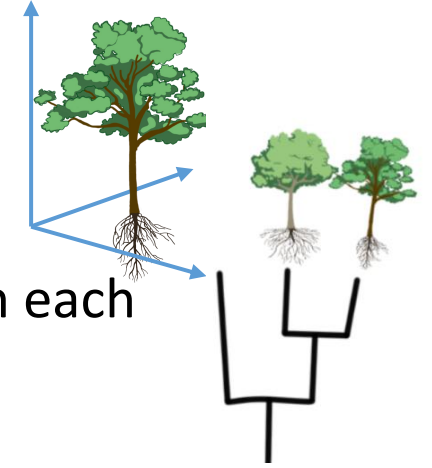
[26] Tucker et al. 2018 [7] Cavender-Bares et al. 2004 [8] Ackerly 2009

H2: Spatial patterns of functional and phylogenetic diversity differ from each other.



H2: Spatial patterns of functional and phylogenetic diversity differ from each other.
→ no clear pattern was found



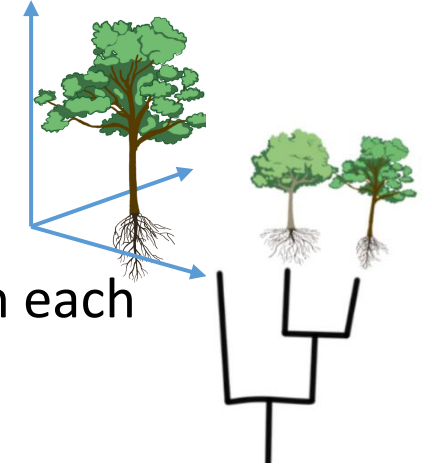


H2: Spatial patterns of functional and phylogenetic diversity differ from each other.

→ no clear pattern was found

Functional diversity differ along rain gradients^[27]

[27] Zuo et al. 2021 [28] Massante et al. 2019 [29] Cai et al. 2020



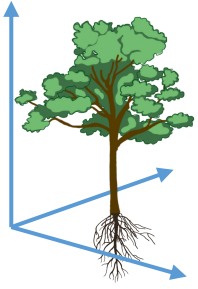
H2: Spatial patterns of functional and phylogenetic diversity differ from each other.
→ no clear pattern was found

Functional diversity differ along rain gradients^[27]

Phylogenetic diversity changes along latitudinal gradient^[28] or to the North (e.g. China)^[29]

[27] Zuo et al. 2021 [28] Massante et al. 2019 [29] Cai et al. 2020

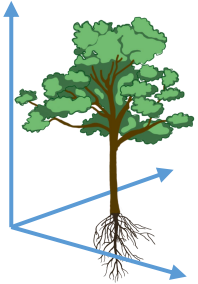
H3: Distribution pattern of functional diversity depends on current climatic conditions.



H3:

Distribution pattern of functional diversity depends on current climatic conditions.

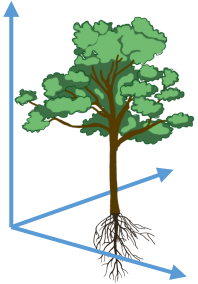
→ explained deviance was relatively low



H3:

Distribution pattern of functional diversity depends on current climatic conditions.

→ explained deviance was relatively low



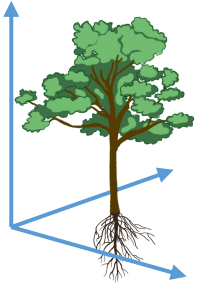
Functional diversity can be linked to recent climate conditions on smaller spatial extent ^[30]

[30] Del Toro et al. 2015 [31] Flynn et al. 2009 [32] Pauw et al. 2021

H3:

Distribution pattern of functional diversity depends on current climatic conditions.

→ explained deviance was relatively low



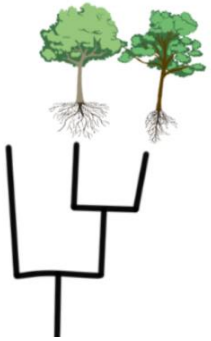
Functional diversity can be linked to recent climate conditions on smaller spatial extent ^[30]

But local communities depend on local factors such as land-use^[31] or soil properties^[32]

[30] Del Toro et al. 2015 [31] Flynn et al. 2009 [32] Pauw et al. 2021

H4:

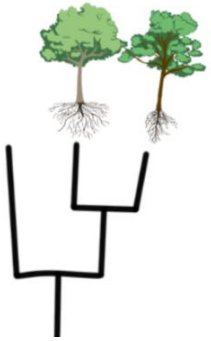
Spatial pattern of phylogenetic diversity depends on past climatic events, i.e. climatic conditions after the last glacial maximum.



H4:

Spatial pattern of phylogenetic diversity depends on past climatic events, i.e. climatic conditions after the last glacial maximum.

→ second most relative influence from BRT, negative correlation in the GAM

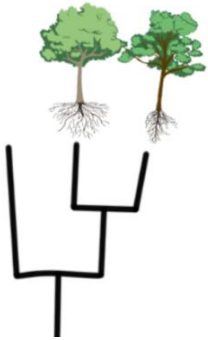


H4:

Spatial pattern of phylogenetic diversity depends on past climatic events, i.e. climatic conditions after the last glacial maximum.

→ second most relative influence from BRT, negative correlation in the GAM

Phylogenetic turnovers in regions with high climatic changes after the LGM^[33]



Abundance weighted indices could improve the understanding of the distribution patterns



05/09/2022

Abundance weighted indices could improve the understanding of the distribution patterns

Addition of local factors could improve model explanation



05/09/2022

Abundance weighted indices could improve the understanding of the distribution patterns

Addition of local factors could improve model explanation

Vegetation-plots from the global South could lead to a better understanding of the observed patterns



Acknowledgements

Thanks to:

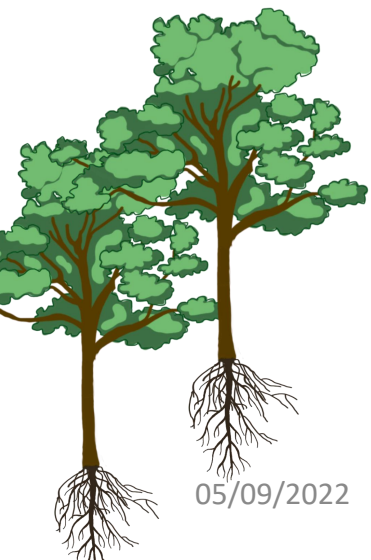
Helge Bruelheide

Francesco M. Sabatini

the IT of the iDiv and the UFZ

the whole working group at the botanical garden

my family and friends.



05/09/2022

Georg Hähn

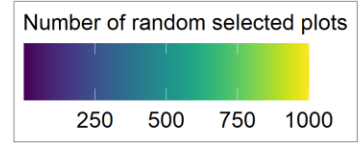
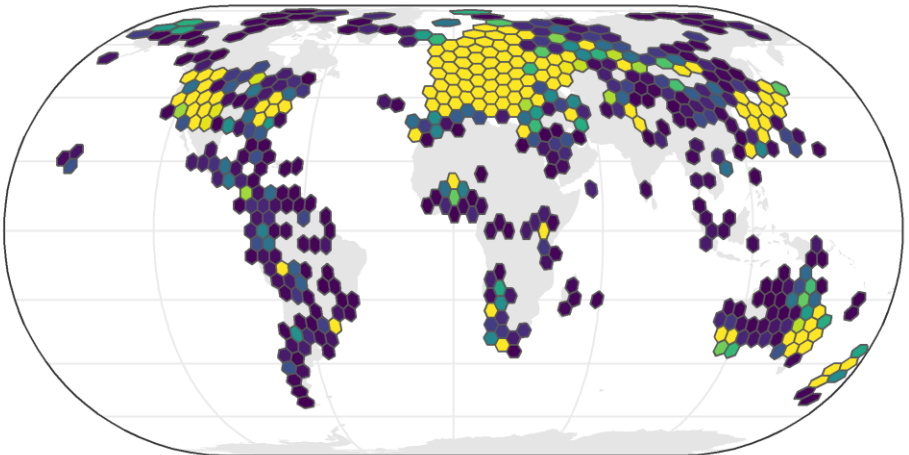
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References

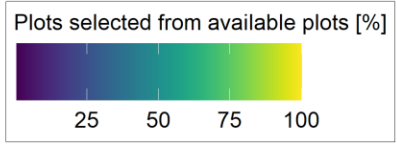
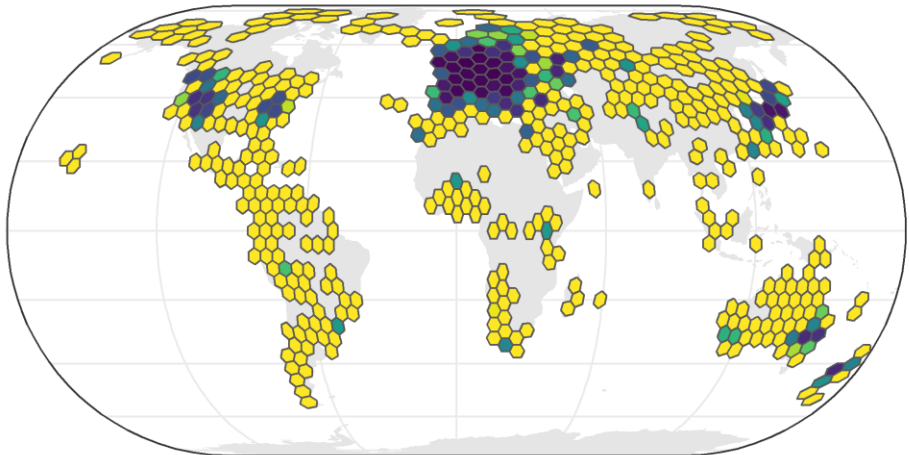
- [1] **Hutchinson**, G. E. (1978), *Introduction to population ecology* (Yale University Press) <<https://agris.fao.org/agris-search/search.do?recordid=us201300539753>>.
- [2] **Gallego-Tévar**, B., Curado, G., Grewell, B. J. et al. (2018), 'Realized niche and spatial pattern of native and exotic halophyte hybrids', *Oecologia*, 188/3: 849–862 <<https://link.springer.com/article/10.1007/s00442-018-4251-y>>.
- [3] **Reich**, P. B., Wright, I. J., Cavender-Bares, J. et al. (2003), 'The Evolution of Plant Functional Variation: Traits, Spectra, and Strategies', *International Journal of Plant Sciences*, 164/S3: S143-S164.
- [4] **García-Palacios**, P., Maestre, F. T., Bardgett, R. D. et al. (2012), 'Plant responses to soil heterogeneity and global environmental change', *J Ecology*, 100/6: 1303-1314.
- [5] **McGill**, B. J., Enquist, B. J., Weiher, E. et al. (2006), 'Rebuilding community ecology from functional traits', *Trends in Ecology & Evolution*, 21/4: 178–185 <<https://www.sciencedirect.com/science/article/pii/S0169534706000334>>.
- [6] **Wright**, I. J., Reich, P. B. and Westoby, M. et al. (2004), 'The worldwide leaf economics spectrum', *Nature*, 428/6985: 821–827 <<https://www.nature.com/articles/nature02403>>.
- [7] **Cavender-Bares**, J., Ackerly, D. D., Baum, D. A. et al. (2004), 'Phylogenetic overdispersion in Floridian oak communities', *The American naturalist*, 163/6: 823–843.
- [8] **Ackerly**, D. (2009), 'Conservatism and diversification of plant functional traits: Evolutionary rates versus phylogenetic signal', *Proceedings of the National Academy of Sciences*, 106 Suppl 2: 19699–19706.
- [9] **Núñez-Farfán**, J., Fornoni, J., and Valverde, P. L. (2007), 'The Evolution of Resistance and Tolerance to Herbivores', *Annual Review of Ecology, Evolution, and Systematics*, 38: 541–566 <<http://www.jstor.org/stable/30033871>>.
- [10] **Castagneyrol**, B., Jactel, H., Vacher, C. et al. (2014), 'Effects of plant phylogenetic diversity on herbivory depend on herbivore specialization', *J Appl Ecol*, 51/1: 134–141 <<https://besjournals.onlinelibrary.wiley.com/doi/full/10.1111/1365-2664.12175>>.
- [11] **Bruelheide**, H., Dengler, J., Jiménez-Alfaro, B. et al. (2019), 'sPlot – A new tool for global vegetation analyses', *J Veg Sci*, 30/2: 161–186 <<https://onlinelibrary.wiley.com/doi/full/10.1111/jvs.12710>>.
- [12] **Smith**, S. A. and **Brown**, J. W. (2018), 'Constructing a broadly inclusive seed plant phylogeny.', *American Journal of Botany*, 105(3): 302–314.
- [13] **Zanne**, A. E., Tank, D. C. and Cornwell, W. K. et al. (2014), 'Three keys to the radiation of angiosperms into freezing environments', *Nature*, 506/7486: 89–92 <<https://www.nature.com/articles/nature12872>>.
- [14] **Shan**, H., Kattge, J. and Reich, P. et al. (2012), 'Gap Filling in the Plant Kingdom - Trait Prediction Using Hierarchical Probabilistic Matrix Factorization', undefined, 2012
- [15] **Fazayeli**, F., Banerjee, A., Kattge, J. et al. (2014), 'Uncertainty Quantified Matrix Completion Using Bayesian Hierarchical Matrix Factorization', *Proceedings - 2014 13th International Conference on Machine Learning and Applications, ICMLA 2014*, 2014: 312–317 <<https://experts.umn.edu/en/publications/uncertainty-quantified-matrix-completion-using-bayesian-hierarchi-2>>.
- [16] **Schrodt**, F., Kattge, J. and Shan, H. et al. (2015), 'BHPMF - a hierarchical Bayesian approach to gap-filling and trait prediction for macroecology and functional biogeography', *Global Ecol Biogeogr*, 24/12: 1510–1521 <<https://onlinelibrary.wiley.com/doi/full/10.1111/geb.12335>>.
- [17] **Kattge**, J., Bönišch, G., Díaz, S. et al. (2020), 'TRY plant trait database - enhanced coverage and open access', *Global Change Biology*, 26/1: 119–188 <<https://onlinelibrary.wiley.com/doi/full/10.1111/gcb.14904>>.

- [18] **Rao**, C. R. (1982), 'Diversity: Its measurement, decomposition, apportionment and analysis', *The Indian Journal of Statistics*, 41: 1–23 <<https://www.jstor.org/stable/25050293>>.
- [19] **Anderson**, M. J., Ellingsen, K. E., and McArdle, B. H. (2006), 'Multivariate dispersion as a measure of beta diversity', *Ecology Letters*, 9/6: 683–693 <https://onlinelibrary.wiley.com/doi/full/10.1111/j.1461-0248.2006.00926.x?casa_token=TRNmHT24BwgAAAAA%3A82ctL876Lf0DfHfsRXqJaVdHHj_COuohj1-yTq8zDPRcNPTB8WKwe8WZVaki8YUk5DEObL-F6A6YOwiv>.
- [20] **Webb**, C. O., Ackerly, D. D. and McPeck, M. A. et al. (2002), 'Phylogenies and Community Ecology', *Annual Review of Ecology and Systematics*, 33/1: 475–505.
- [21] **Botta-Dukát**, Z. (2018), 'Cautionary note on calculating standardized effect size (SES) in randomization test', *Community Ecology*, 19/1: 77–83 <<https://akjournals.com/view/journals/168/19/1/article-p77.xml>>.
- [22] **Karger**, D. N., Conrad, O., Böhner, J. et al. (2017), 'Climatologies at high resolution for the earth's land surface areas', *Sci Data*, 4: 170122.
- [23] **Karger**, D. N., Conrad, O., Böhner, J. et al. (2018), *Data from: Climatologies at high resolution for the earth's land surface areas*.
- [24] **Brown**, S. C., Wigley, T. M. L., Otto-Bliesner, B. L. et al. (2020), 'StableClim, continuous projections of climate stability from 21000 BP to 2100 CE at multiple spatial scales', *Sci Data*, 7/1: 335 <<https://www.nature.com/articles/s41597-020-00663-3#Sec20>>.
- [25] **Colin**, B., Schmidt, M., Clifford, S. et al. (2018), 'Influence of Spatial Aggregation on Prediction Accuracy of Green Vegetation Using Boosted Regression Trees', *Remote Sens*, 10/8.
- [26] **Tucker**, C. M., Davies, T. J. and Cadotte, M. W. et al. (2018), 'On the relationship between phylogenetic diversity and trait diversity', *Ecology*, 99/6: 1473–1479 <https://esajournals.onlinelibrary.wiley.com/doi/full/10.1002/ecy.2349?casa_token=ujZVwD1q-UAAAAA%3AN8GjICFEo8ohMet2kpFEwL9RLzffn7WLTR1cQH4YR33EIN8zQMUHQyZGc6eJ9vA2ZqEDvN44tT7CoHg>.
- [27] **Zuo**, X., Zhao, S. and Cheng, H. et al. (2021), 'Functional diversity response to geographic and experimental precipitation gradients varies with plant community type', *Funct Ecol*, 35/9: 2119–2132 <<https://besjournals.onlinelibrary.wiley.com/doi/full/10.1111/1365-2435.13875>>.
- [28] **Massante**, J. C., Götzenberger, L., Takkis, K. et al. (2019), 'Contrasting latitudinal patterns in phylogenetic diversity between woody and herbaceous communities', *Sci Rep*, 9/1: 6443 <<https://www.nature.com/articles/s41598-019-42827-1>>.
- [29] **Cai**, H., Lyu, L., Shrestha, N. et al. (2021), 'Geographical patterns in phylogenetic diversity of Chinese woody plants and its application for conservation planning', *Divers Distrib*, 27/1: 179–194 <<https://onlinelibrary.wiley.com/doi/full/10.1111/ddi.13180>>.
- [30] **Del Toro**, I., Silva, R. R., and Ellison, A. M. (2015), 'Predicted impacts of climatic change on ant functional diversity and distributions in eastern North American forests', *Diversity Distrib.*, 21/7: 781–791 <<https://onlinelibrary.wiley.com/doi/full/10.1111/ddi.12331>>.
- [31] **Flynn**, D. F. B., Gogol-Prokurat, M., Nogeire, T. et al. (2009), 'Loss of functional diversity under land use intensification across multiple taxa', *Ecology Letters*, 12/1: 22–33 <<https://onlinelibrary.wiley.com/doi/full/10.1111/j.1461-0248.2008.01255.x>>.
- [32] **Pauw**, K. de, Meeussen, C., Govaert, S. et al. (2021), 'Taxonomic, phylogenetic and functional diversity of understorey plants respond differently to environmental conditions in European forest edges', *J Ecology*, 109/7: 2629–2648 <<https://besjournals.onlinelibrary.wiley.com/doi/full/10.1111/1365-2745.13671>>.
- [33] **Cubino**, J. P., Jiménez-Alfaro, B., Sabatini, F. M. et al. (2020), 'Plant taxonomic and phylogenetic turnover increases toward climatic extremes and depends on historical factors in European beech forests', *Journal of Vegetation Science*, 32/1 <<https://onlinelibrary.wiley.com/doi/full/10.1111/jvs.12977>>.

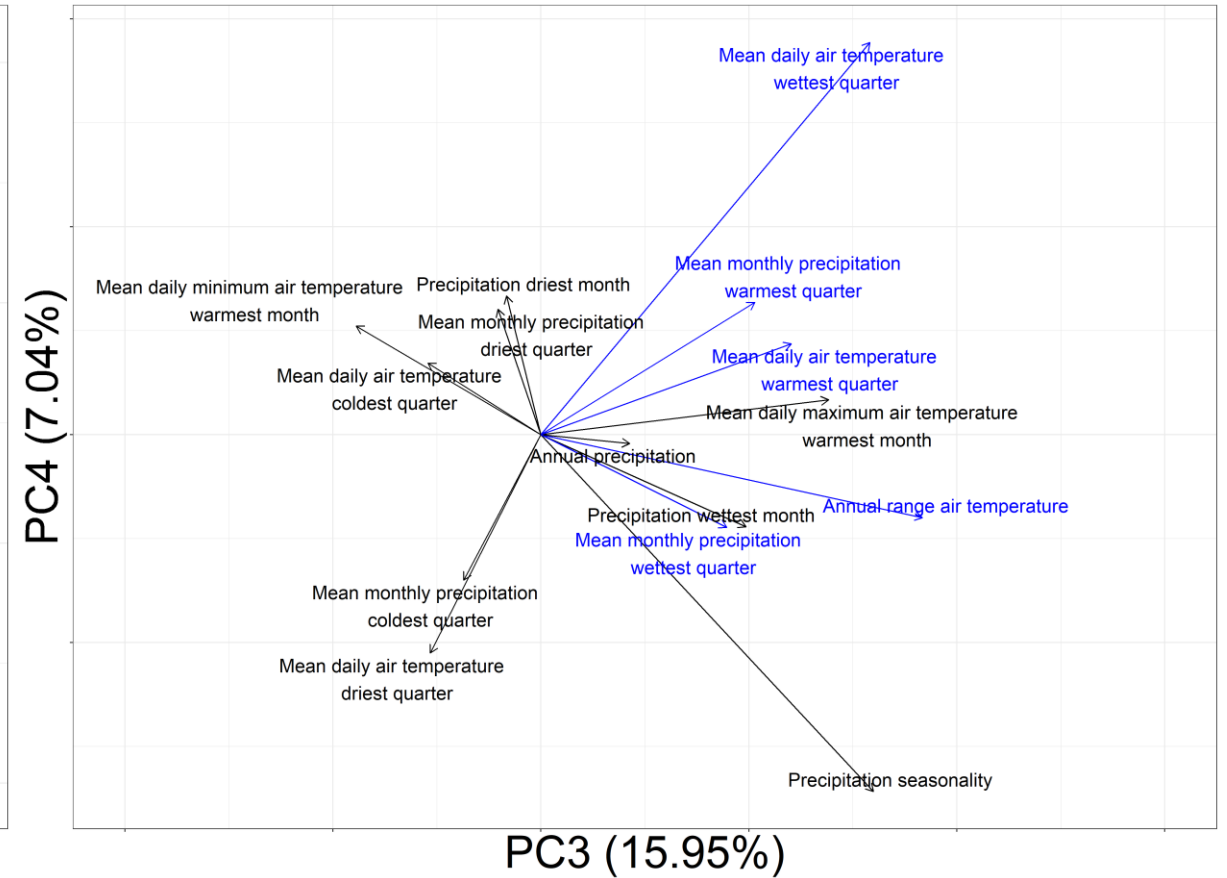
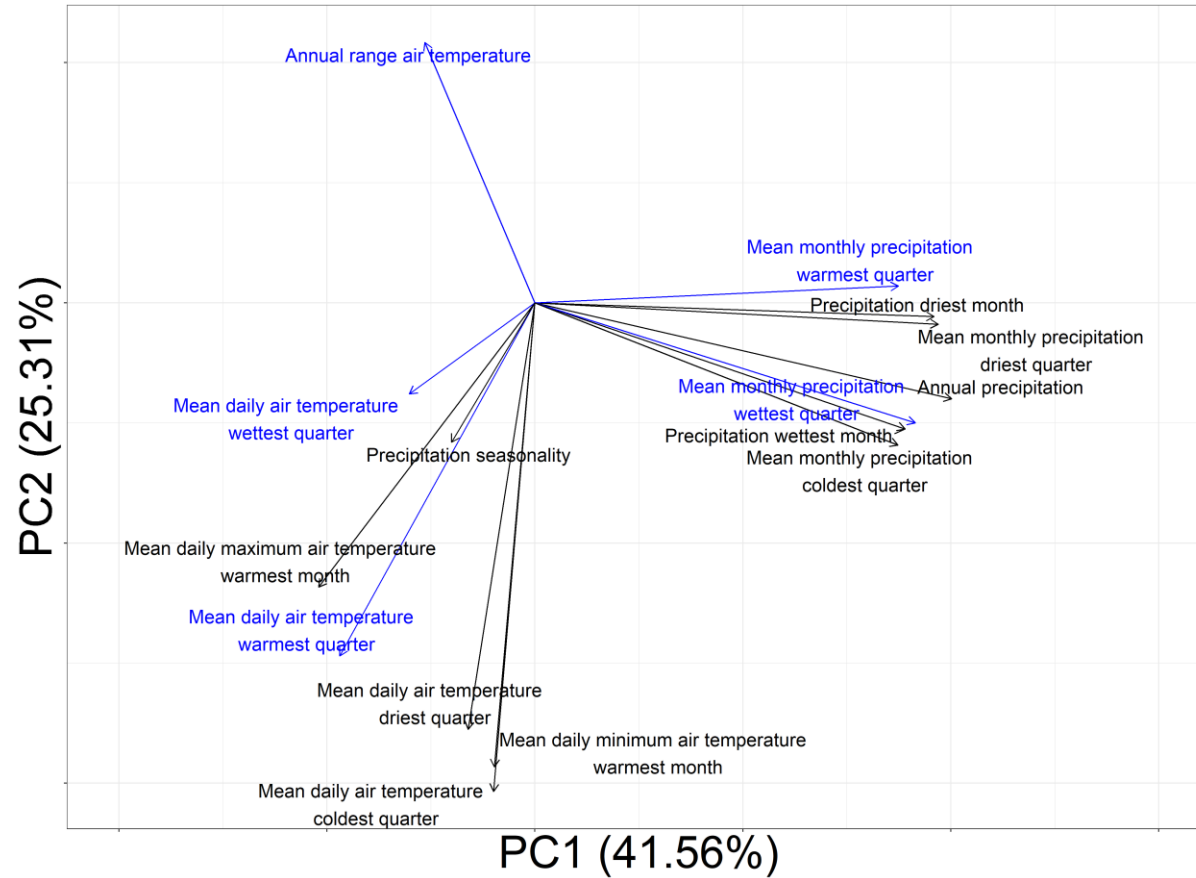
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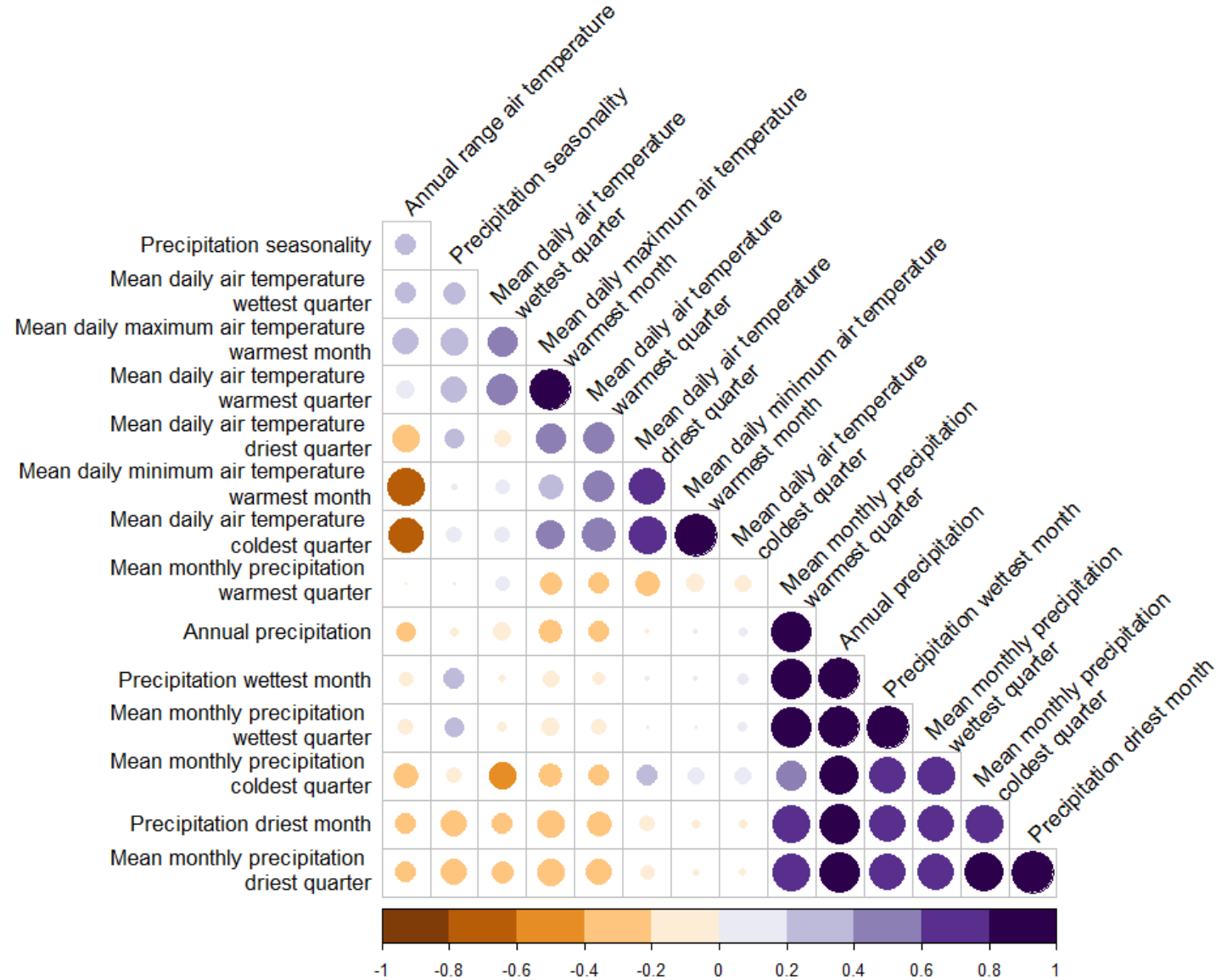
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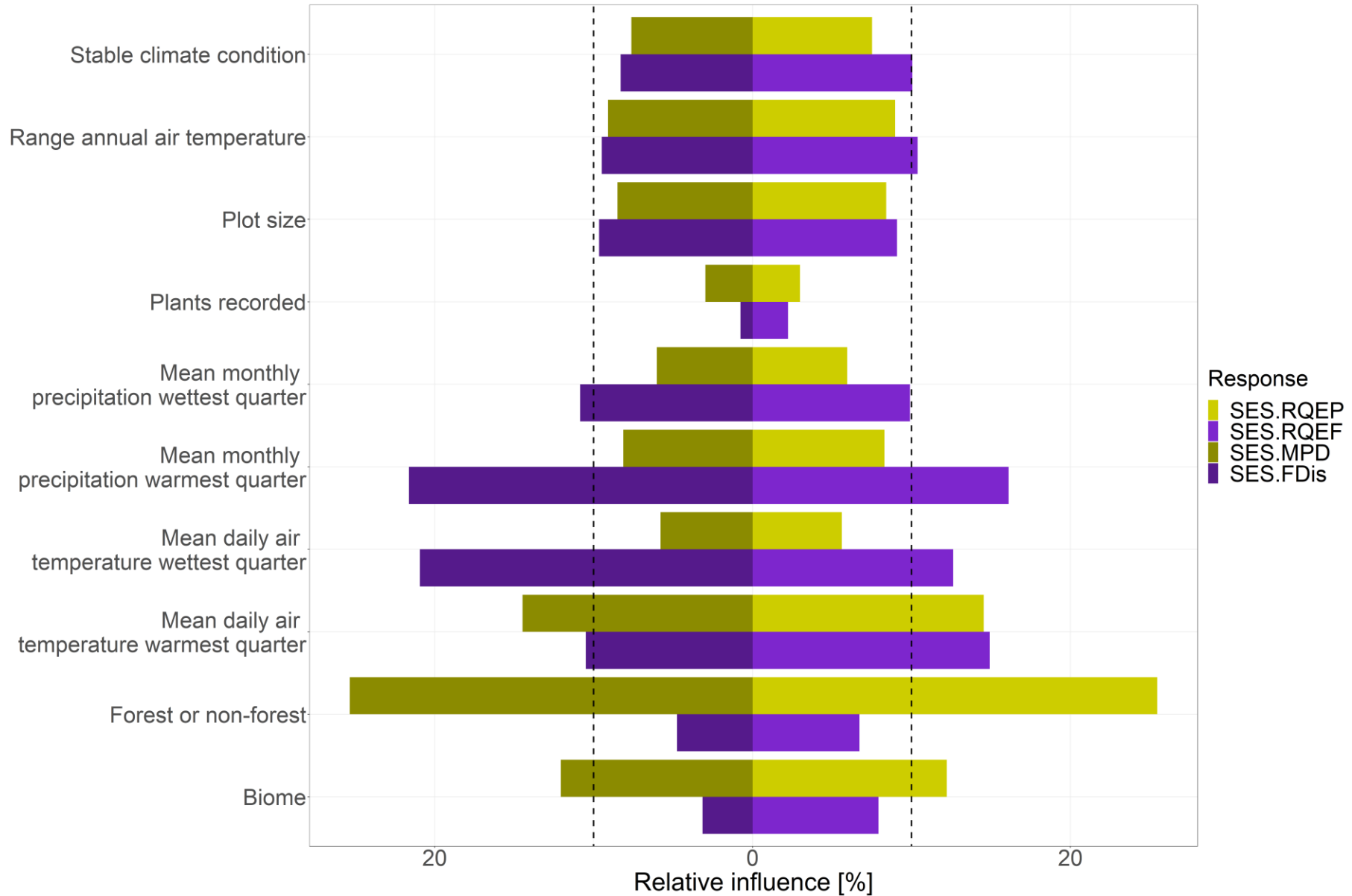
Supporting Information



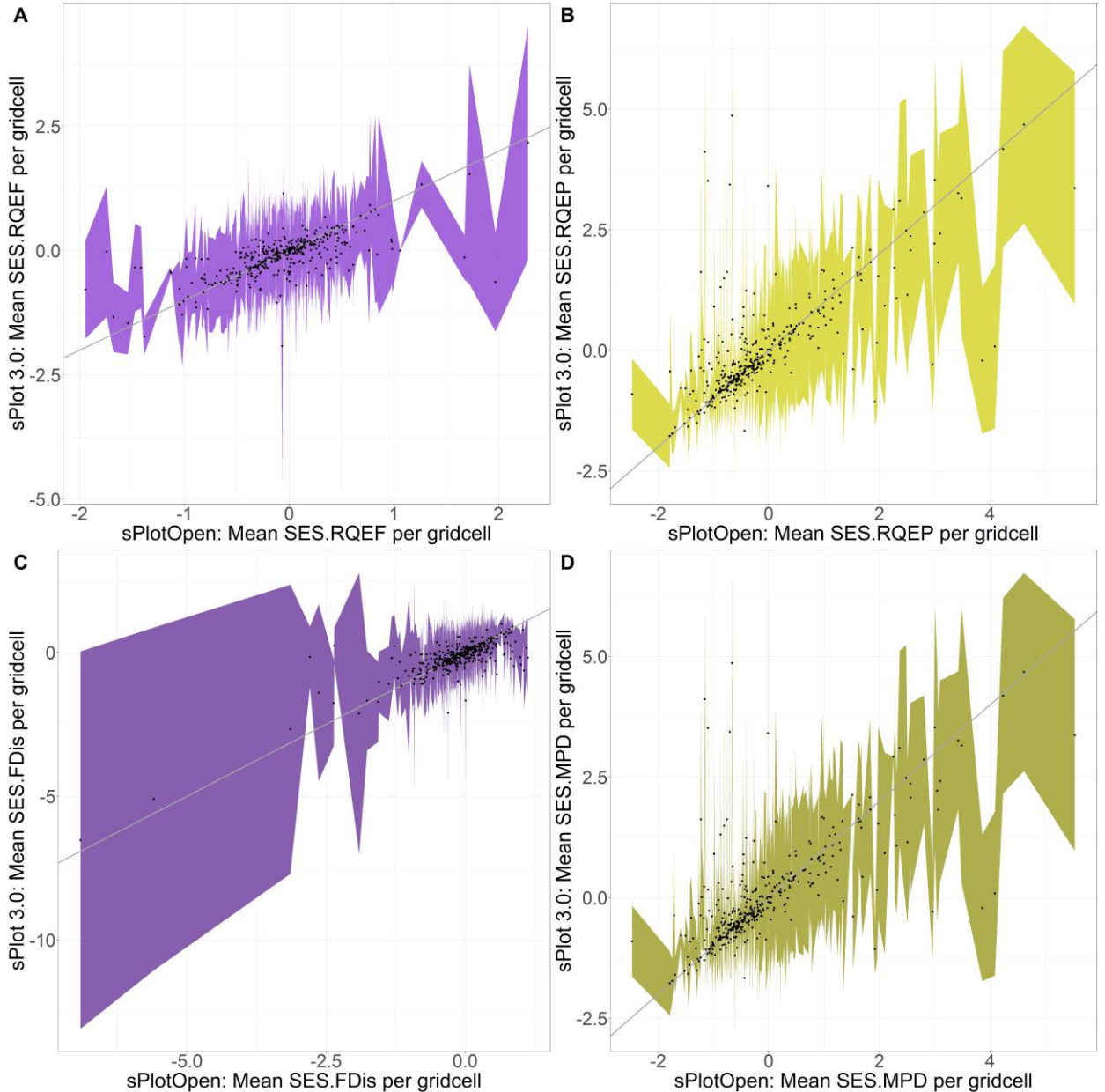
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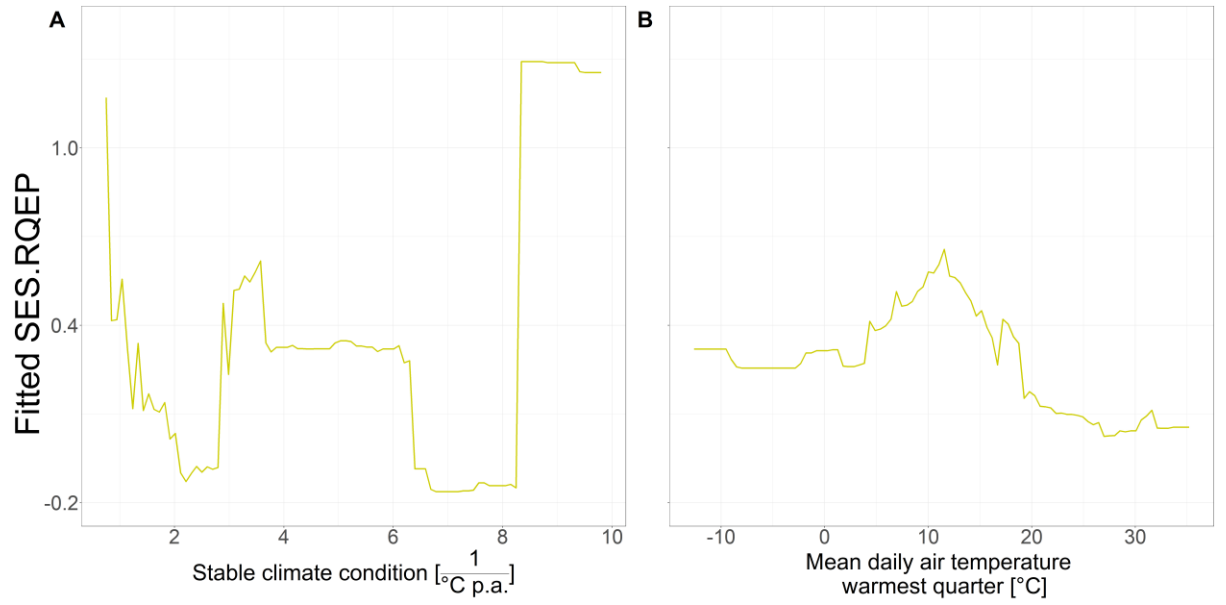
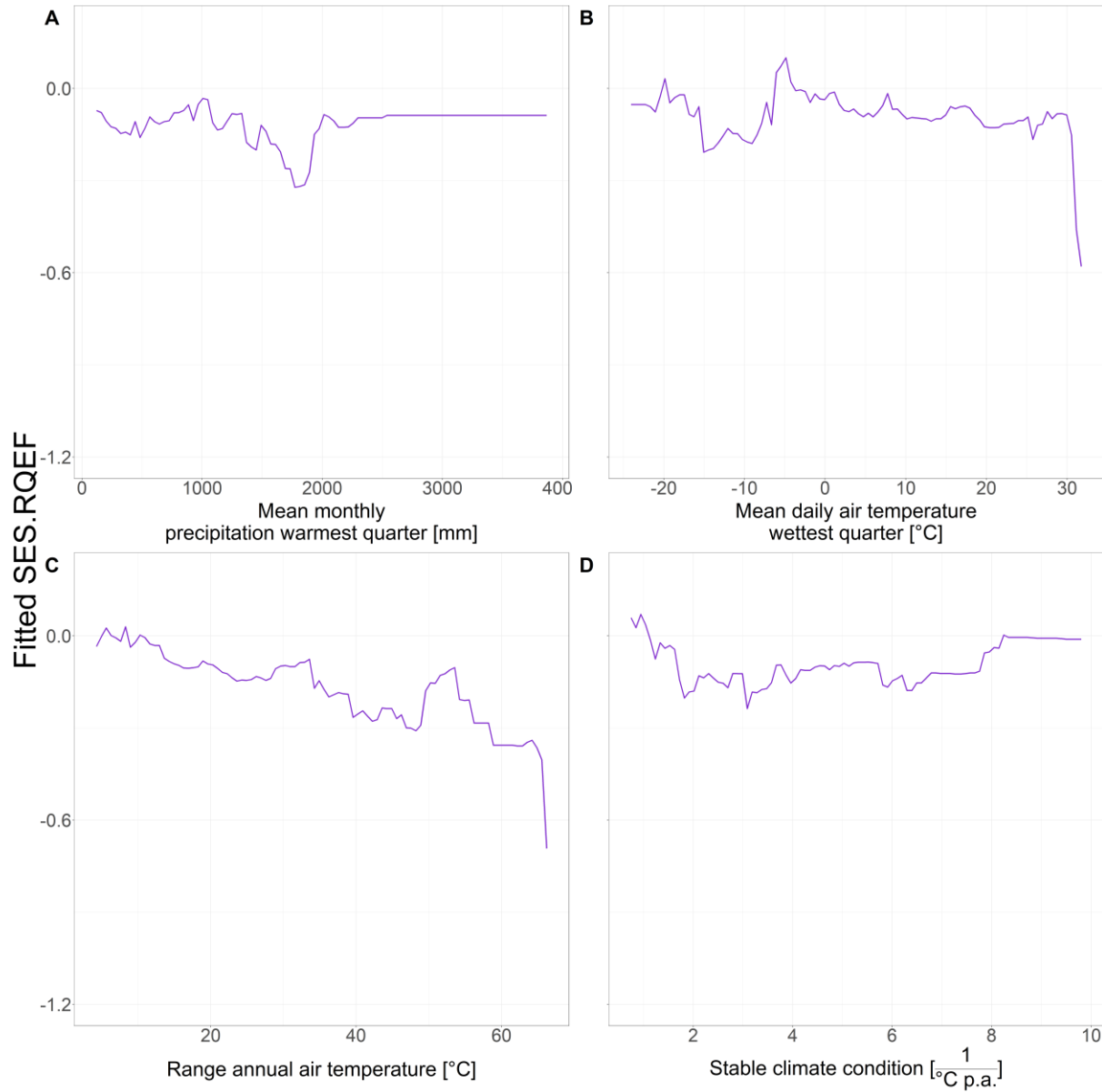
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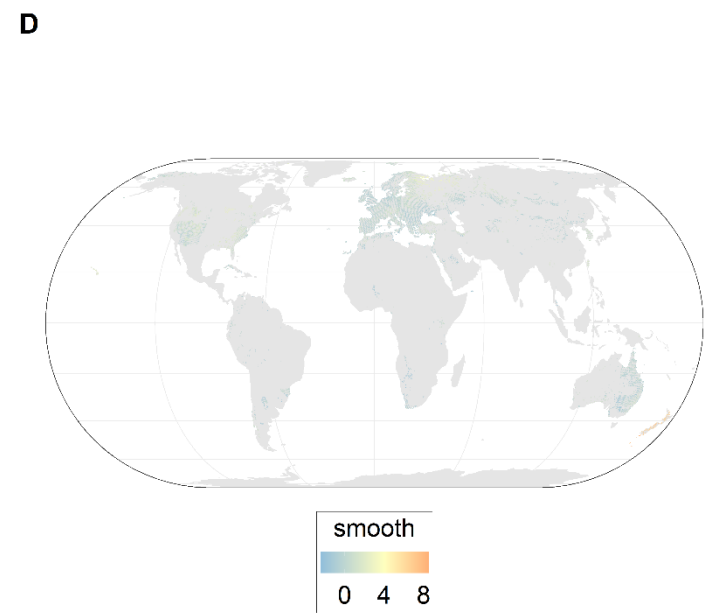
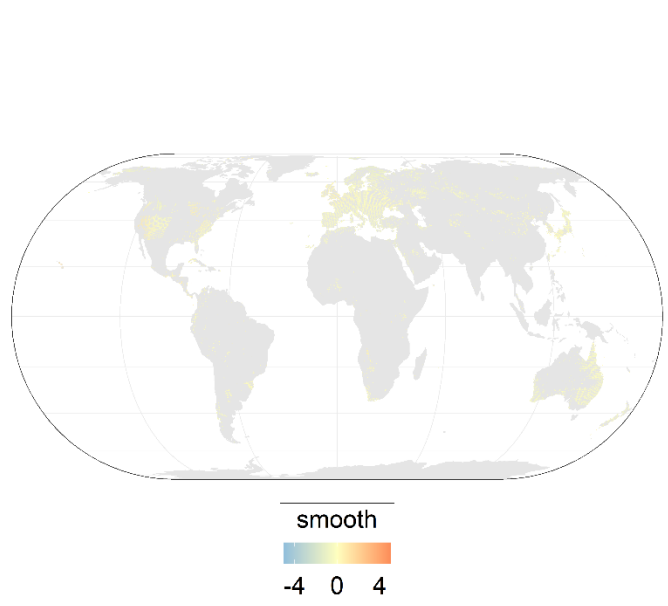
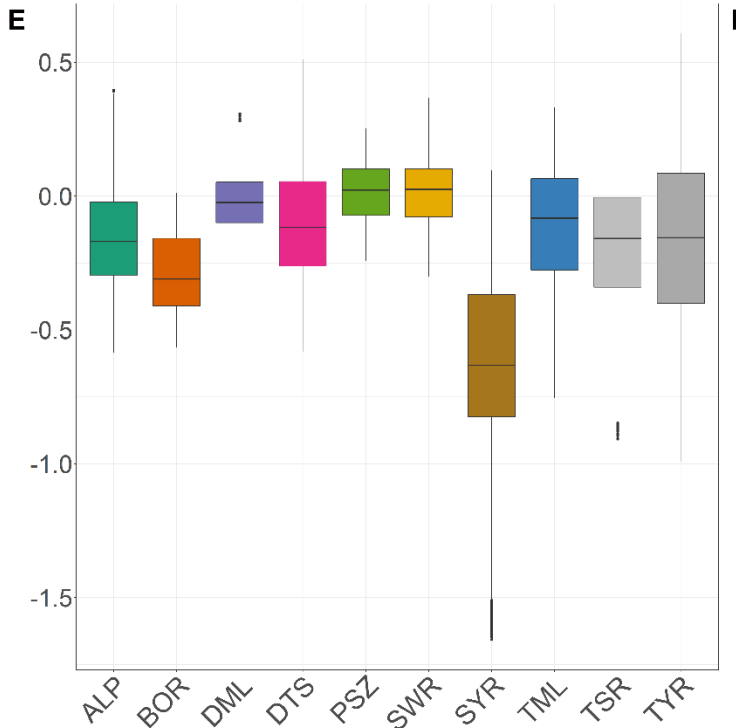
Supporting Information



Supporting Information



Supporting Information



Supporting Information

```
Family: gaussian
Link function: identity

Formula:
SES.RQEF ~ SES.RQEF + s(Longitude, Latitude, bs = "sos")

Parametric coefficients:
      Estimate Std. Error t value Pr(>|t|)
(Intercept)  0.0001157  0.0007246    0.16   0.873
SES.RQEF     -0.0896341  0.0004374  -204.94 <2e-16 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Approximate significance of smooth terms:
      edf Ref.df    F p-value
s(Longitude, Latitude) 48.94    49 1281 <2e-16 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

R-sq.(adj) = 0.0581  Deviance explained = 5.81%
-REML = 2.4684e+06  Scale est. = 0.93336    n = 1782777

Family: gaussian
Link function: identity

Formula:
SES.RQEF ~ stable.clim + mean.daily.air.temp.warm.qu + s(Longitude,
  Latitude, bs = "sos") + is.forest

Parametric coefficients:
      Estimate Std. Error t value Pr(>|t|)
(Intercept)    1.0286811  0.0118456   86.84 <2e-16 ***
stable.clim    -0.1694141  0.0071234  -23.78 <2e-16 ***
mean.daily.air.temp.warm.qu -0.0733673  0.0004541 -161.56 <2e-16 ***
is.forestTRUE    1.8155171  0.0027595   657.91 <2e-16 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Approximate significance of smooth terms:
      edf Ref.df    F p-value
s(Longitude, Latitude) 48.96    49 4802 <2e-16 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

R-sq.(adj) = 0.394  Deviance explained = 39.4%
-REML = 2.696e+06  Scale est. = 2.1407    n = 1498079
```

```
Family: gaussian
Link function: identity

Formula:
SES.RQEF ~ stable.clim + annual.range.air.temp + mean.monthly.prec.warm.qu +
  mean.daily.air.temp.wet.qu + biome + s(Longitude, Latitude,
  bs = "sos")

Parametric coefficients:
      Estimate Std. Error t value Pr(>|t|)
(Intercept)   -4.923e-02  1.296e-02  -3.798 0.000146 ***
stable.clim    6.378e-02  4.201e-03  15.181 < 2e-16 ***
annual.range.air.temp 1.986e-03  3.008e-04   6.604 4.01e-11 ***
mean.monthly.prec.warm.qu -4.423e-04  7.611e-06 -58.106 < 2e-16 ***
mean.daily.air.temp.wet.qu 4.950e-03  1.846e-04  26.813 < 2e-16 ***
biomeBoreal zone -1.867e-01  9.579e-03 -19.485 < 2e-16 ***
biomeDry midlatitudes 4.680e-02  8.611e-03   5.435 5.48e-08 ***
biomeDry tropics and subtropics 4.159e-02  1.098e-02   3.789 0.000151 ***
biomePolar and subpolar zone 1.202e-01  2.021e-02   5.951 2.67e-09 ***
biomeSubtrop. with year-round rain -1.662e-01  1.009e-02 -16.473 < 2e-16 ***
biomeSubtropics with winter rain -3.242e-02  7.556e-03  -4.290 1.79e-05 ***
biomeTemperate midlatitudes -2.699e-02  6.463e-03  -4.176 2.96e-05 ***
biomeTropics with summer rain -1.349e-02  1.671e-02  -0.808 0.419360
biomeTropics with year-round rain -5.631e-03  1.903e-02  -0.296 0.767339
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Approximate significance of smooth terms:
      edf Ref.df    F p-value
s(Longitude, Latitude) 48.93    49 1021 <2e-16 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

R-sq.(adj) = 0.0397  Deviance explained = 3.98%
-REML = 2.4856e+06  Scale est. = 0.95156    n = 1782777
```


Supporting Information

```
R version 4.2.1 (2022-06-23)
Platform: x86_64-pc-linux-gnu (64-bit)
Running under: Ubuntu 20.04.5 LTS

Matrix products: default
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LAPACK: /usr/lib/x86_64-linux-gnu/lapack/liblapack.so.3.9.0

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 [6] LC_MESSAGES=en_US.UTF-8  LC_PAPER=en_US.UTF-8     LC_NAME=C                LC_ADDRESS=C            LC_TELEPHONE=C
[11] LC_MEASUREMENT=en_US.UTF-8 LC_IDENTIFICATION=C

attached base packages:
[1] stats      graphics  grDevices  utils      datasets  methods    base

other attached packages:
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