

# The distribution patterns of functional and phylogenetic diversity in vascular plant communities.

Plot

Edinburgh, Scotland

 $18^{\text{th}} - 21^{\text{st}}$  December 2022

• iDiv

<u>Georg J. A. Hähn</u>, Francesco M. Sabatini, Gabriella Damasceno, and Helge Bruelheide

> 21/12/2022 12:15:00 Fintry Auditorium



**BES 2022** 

Annual Meeting





BES 2022 Annual Meeting

Introduction – Plant species niche







Introduction – Plant species niche in the environment





**BES 2022** 

Annual Meeting





BES 2022 Annual Meeting

Introduction – Analysis of the niche







Introduction – Analysis the niche: Traits



**BES 2022** 

Annual Meeting





BES 2022 Annual Meeting

Introduction – Analysis of the niche: Traits







BES 2022 Annual Meeting

Introduction – Analysis of the niche: Traits

Specific leaf area (SLA) low → species tolerates stress high → most use of favourable conditions

Specific root length (**SRL**) → nutrient and water uptake







Introduction – Analysis of the niche: Phylogeny



**BES 2022** 

Annual Meeting







**BES 2022** Annual Meeting

Introduction – Analysis of the niche: Phylogeny









BES 2022 Annual Meeting

Introduction – Analysis of the niche: Phylogeny







MART HALL

MARTIN-LUTHER-UNIVERSITÄT Georg J. A. Hähn HALLE-WITTENBERG BES 2022 Annual Meeting

Introduction – Analysis of the niche: Phylogeny





[3] Reich et al. 2003 [6] Cavender-Bares et al. 2004 [7] Ackerly 2009





BES 2022 Annual Meeting

Introduction – Research questions

1. Are functional and phylogenetic diversity correlated with each other at the global scale?





Plot

MARTIN-LUTHER-UNIVERSITÄT Georg J. A. Hähn HALLE-WITTENBERG BES 2022

Annual Meeting

Introduction – Research questions

1. Are functional and phylogenetic diversity correlated with each other at the global scale?

2. To what extent is the geographical distribution of functional and phylogenetic diversity explained by present and past climate conditions?



Methods – Response and explanatory variables

• iDiv

Edinburgh, Scotland

18<sup>th</sup> – 21<sup>st</sup> December 2022



a project of 💿 iDiv

**BES 2022** 

Annual Meeting

Methods – Response and explanatory variables

• iDiv

Edinburgh, Scotland

18<sup>th</sup> – 21<sup>st</sup> December 2022



Plot

**BES 2022** 

Annual Meeting

Plot MARTI

16

MARTIN-LUTHER-UNIVERSITÄT Georg J. A. Hähn HALLE-WITTENBERG **BES 2022** 

Annual Meeting

Methods – Response and explanatory variables

iDiv

Edinburgh, Scotland

 $18^{\text{th}} - 21^{\text{st}}$  December 2022







MARTIN-LUTHER-UNIVERSITÄT Georg J. A. Hähn HALLE-WITTENBERG BES 2022 Annual Meeting

## Methods – Response and explanatory variables



[13] Smith and Brown 2018 [14] Zanne et al. 2014





MARTIN-LUTHER-UNIVERSITÄT Georg J. A. Hähn HALLE-WITTENBERG BES 2022 Annual Meeting



## iDiv 🕝 sPlot



MARTIN-LUTHER-UNIVERSITÄT Georg J. A. Hähn HALLE-WITTENBERG BES 2022 Annual Meeting





Plot

MARTIN-LUTHER-UNIVERSITÄT Georg J. A. Hähn HALLE-WITTENBERG

BES 2022 Annual Meeting



## iDiv 🕘 sPlot



MARTIN-LUTHER-UNIVERSITÄT Georg J. A. Hähn HALLE-WITTENBERG BES 2022 Annual Meeting





MARTIN-LUTHER-UNIVERSITÄT Georg J. A. Hähn HALLE-WITTENBERG

BES 2022 Annual Meeting

## Methods – Response and explanatory variables



iDiv

## iDiv 🕘 sPlot



MARTIN-LUTHER-UNIVERSITÄT Georg J. A. Hähn HALLE-WITTENBERG BES 2022 Annual Meeting



Edinburgh, Scotland  $18^{\text{th}} - 21^{\text{st}}$  December 2022





**BES 2022** Annual Meeting

## Methods – Response and explanatory variables

iDiv



[15] Rao 1982

#### Edinburgh, Scotland **BES 2022** iDiv Plot MARTIN-LUTHER-UNIVERSITÄT Georg J. A. Hähn $18^{\text{th}} - 21^{\text{st}}$ December 2022 **Annual Meeting** HALLE-WITTENBERG Methods – Response and explanatory variables Predictors Specific leaf area, plant height **SPlot** and specific root length from gap-filled TRY traits<sup>[9, 10, 11, 12]</sup> a project of ( ) iDiv **Plant Trait Database** Plot size GBOTB seed plants<sup>[13]</sup> Forest or non-forest Clade in the phylogeny Presence-absence data for Biome<sup>[8]</sup> for pteridophytes<sup>[14]</sup> 1,977,637 vegetation-plots<sup>[8]</sup> 50% of the total cover is represented by species for which phylogenetic and trait data were available →1,782,777 plots Phylogenetic diversity: Functional diversity: Rao`s Quadratic Entropy<sup>[15]</sup> Rao`s Quadratic Entropy<sup>[15]</sup> $\sum^{n-1} \sum^n d_{ij} p_i p_j$ **gam**( Functional Entropy ~ Phylogenetic Entropy + s(Longitude, Latitude, RQE =bs = "sos"), family = "gaussian", method = "REML") $\_$ <u>observed - expected</u> SES

[8] Bruelheide et al. 2019

 $\overline{SD}_{expected}$ 

#### Edinburgh, Scotland **BES 2022** iDiv Plot MARTIN-LUTHER-UNIVERSITÄT Georg J. A. Hähn $18^{\text{th}} - 21^{\text{st}}$ December 2022 Annual Meeting HALLE-WITTENBERG Predictors Methods – Response and explanatory variables Specific leaf area, plant height **SPlot CHFLSA** and specific root length from gap-filled TRY traits<sup>[9, 10, 11, 12]</sup> 19 bioclimatic variables Plant Trait Database a project of ( **b i Div** from CHELSA v.2.1<sup>[16, 17]</sup> Plot size GBOTB seed plants<sup>[13]</sup> Forest or non-forest Clade in the phylogeny Presence-absence data for Biome<sup>[8]</sup> for pteridophytes<sup>[14]</sup> 1,977,637 vegetation-plots<sup>[8]</sup> 50% of the total cover is represented by species for which phylogenetic and trait data were available →1,782,777 plots Phylogenetic diversity: Functional diversity: Rao`s Quadratic Entropy<sup>[15]</sup> Rao`s Quadratic Entropy<sup>[15]</sup> **gam**( Functional Entropy ~ Phylogenetic Entropy + s(Longitude, Latitude, $\sum_{ij}^{n} \sum_{j=1}^{n} d_{ij}p_ip_j$ RQE =bs = "sos"), family = "gaussian", method = "REML") $\_$ <u>observed - expected</u> SES $\overline{SD}_{expected}$

26

#### Edinburgh, Scotland **BES 2022** iDiv Plot MARTIN-LUTHER-UNIVERSITÄT Georg J. A. Hähn $18^{\text{th}} - 21^{\text{st}}$ December 2022 Annual Meeting HALLE-WITTENBERG Predictors Methods – Response and explanatory variables TraCE-21ka (paleo climate) Specific leaf area, plant height CHFLSA **SPlot** and specific root length from gap-filled TRY traits<sup>[9, 10, 11, 12]</sup> 19 bioclimatic variables a project of ( **b i Div** Worldwide stable **Plant Trait Database** from CHELSA v.2.1<sup>[16, 17]</sup> climatic condition after Plot size GBOTB seed plants<sup>[13]</sup> last glacial maximum Forest or non-forest Clade in the phylogeny Presence-absence data for (LGM) from StableClim Biome<sup>[8]</sup> for pteridophytes<sup>[14]</sup> 1,977,637 vegetation-plots<sup>[8]</sup> $v.1.1^{[18]}$ 50% of the total cover is represented by species for which phylogenetic and trait data were available →1,782,777 plots Phylogenetic diversity: Functional diversity: Rao`s Quadratic Entropy<sup>[15]</sup> Rao`s Quadratic Entropy<sup>[15]</sup> $\sum^{n-1} \sum^{n} d_{ij} p_i p_j$ **gam**( Functional Entropy ~ Phylogenetic Entropy + s(Longitude, Latitude, RQE =bs = "sos"), family = "gaussian", method = "REML") $\_$ <u>observed - expected</u> SES $SD_{expected}$

[18] Brown et al. 2020

#### Edinburgh, Scotland **BES 2022** iDiv Plot MARTIN-LUTHER-UNIVERSITÄT Georg J. A. Hähn $18^{\text{th}} - 21^{\text{st}}$ December 2022 Annual Meeting HALLE-WITTENBERG Predictors Methods – Response and explanatory variables TraCE-21ka (paleo climate) Specific leaf area, plant height **CHFLSA SPlot** and specific root length from gap-filled TRY traits<sup>[9, 10, 11, 12]</sup> 19 bioclimatic variables a project of ( iDiv Worldwide stable **Plant Trait Database** from CHELSA v.2.1<sup>[16, 17]</sup> climatic condition after Plot size GBOTB seed plants<sup>[13]</sup> last glacial maximum Forest or non-forest Clade in the phylogeny Presence-absence data for (LGM) from StableClim Biome<sup>[8]</sup> for pteridophytes<sup>[14]</sup> 1,977,637 vegetation-plots<sup>[8]</sup> $v.1.1^{[18]}$ 50% of the total cover is represented by **Principial component** species for which phylogenetic and analyses (PCA) & trait data were available **boosted regression** trees →1,782,777 plots Phylogenetic diversity: Functional diversity: Rao`s Quadratic Entropy<sup>[15]</sup> Rao`s Quadratic Entropy<sup>[15]</sup> $\sum_{i=1}^{n-1}\sum_{j=1}^{n}d_{ij}p_ip_j$ **gam**( Functional Entropy ~ Phylogenetic Entropy + s(Longitude, Latitude, RQE =bs = "sos"), family = "gaussian", method = "REML") $\_$ <u>observed - expected</u> SES : SD<sub>expected</sub>

28

[18] Brown et al. 2020

#### Edinburgh, Scotland **BES 2022** iDiv Plot MARTIN-LUTHER-UNIVERSITÄT Georg J. A. Hähn $18^{\text{th}} - 21^{\text{st}}$ December 2022 Annual Meeting HALLE-WITTENBERG Methods – Response and explanatory variables Predictors TraCE-21ka (paleo climate) Specific leaf area, plant height **CHFLSA SPlot** and specific root length from gap-filled TRY traits<sup>[9, 10, 11, 12]</sup> 19 bioclimatic variables a project of 🌘 iDiv Worldwide stable **Plant Trait Database** from CHELSA v.2.1<sup>[16, 17]</sup> climatic condition after Plot size GBOTB seed plants<sup>[13]</sup> last glacial maximum Forest or non-forest Clade in the phylogeny Presence-absence data for (LGM) from StableClim Biome<sup>[8]</sup> for pteridophytes<sup>[14]</sup> 1,977,637 vegetation-plots<sup>[8]</sup> $v.1.1^{[18]}$ 50% of the total cover is represented by **Principial component** species for which phylogenetic and analyses (PCA) & trait data were available **boosted** regression trees $\rightarrow$ 5 recent climate variables →1,782,777 plots $\rightarrow$ climate variability after LGM Phylogenetic diversity: $\rightarrow$ 3 plot variables Functional diversity: Rao`s Quadratic Entropy<sup>[15]</sup> Rao`s Quadratic Entropy<sup>[15]</sup> $\overline{\boldsymbol{\mathcal{S}}}^{n-1}$ $\overline{\boldsymbol{\mathcal{S}}}^{n-1}$ $d_{ij}p_ip_j$ **gam**( Functional Entropy ~ Phylogenetic Entropy + s(Longitude, Latitude, RQE =bs = "sos"), family = "gaussian", method = "REML") $\_$ observed – expected SES :

 $SD_{expected}$ 

#### Edinburgh, Scotland **BES 2022** iDiv Plot MARTIN-LUTHER-UNIVERSITÄT Georg J. A. Hähn $18^{\text{th}} - 21^{\text{st}}$ December 2022 Annual Meeting HALLE-WITTENBERG Methods – Response and explanatory variables Predictors TraCE-21ka (paleo climate) Specific leaf area, plant height **CHFLSA SPlot** and specific root length from gap-filled TRY traits<sup>[9, 10, 11, 12]</sup> 19 bioclimatic variables a project of 🌘 iDiv Worldwide stable **Plant Trait Database** from CHELSA v.2.1<sup>[16, 17]</sup> climatic condition after Plot size GBOTB seed plants<sup>[13]</sup> last glacial maximum Forest or non-forest Clade in the phylogeny Presence-absence data for (LGM) from StableClim Biome<sup>[8]</sup> for pteridophytes<sup>[14]</sup> 1,977,637 vegetation-plots<sup>[8]</sup> $v.1.1^{[18]}$ 50% of the total cover is represented by **Principial component** species for which phylogenetic and analyses (PCA) & trait data were available **boosted** regression trees $\rightarrow$ 5 recent climate variables →1,782,777 plots $\rightarrow$ climate variability after LGM Phylogenetic diversity: $\rightarrow$ 3 plot variables Functional diversity: Rao`s Quadratic Entropy<sup>[15]</sup> Rao`s Quadratic Entropy<sup>[15]</sup> **gam**( Functional Entropy ~ Phylogenetic Entropy + s(Longitude, Latitude, RQE = $\langle n^{n-1} \rangle = d_{ij}p_ip_j$ bs = "sos"), family = "gaussian", method = "REML") **gam**( Entropy ~ Explanatory variables + s(Longitude, Latitude, \_ observed – expected *bs* = "sos"), *family* = "gaussian", *method* = "REML") SES : $SD_{expected}$





МА НА

MARTIN-LUTHER-UNIVERSITÄT Georg J. A. Hähn HALLE-WITTENBERG

BES 2022 Annual Meeting

Results – Distribution of functional and phylogenetic entropy





MARTIN-LUTHER-UNIVERSITÄT Georg J. A. Hähn HALLE-WITTENBERG **BES 2022** 

**Annual Meeting** 

### Results – Relationship of functional and phylogenetic entropy

iDiv

**gam**(Functional Entropy ~ Phylogenetic Entropy + s(Longitude, Latitude, bs = "sos"), family = "gaussian", method = "REML")

Plot



coldest quarter [°C]

## Plot 🚳

MARTIN-LUTHER-UNIVERSITÄT Georg J. A. Hähn HALLE-WITTENBERG

### BES 2022 Annual Meeting













Discussion – Are functional and phylogenetic diversity correlated at the global scale?

Functional and phylogenetic diversity are negatively correlated at the global scale





BES 2022 Annual Meeting

Discussion – Are functional and phylogenetic diversity correlated at the global scale?

Functional and phylogenetic diversity are negatively correlated at the global scale







BES 2022 Annual Meeting

Discussion – Are functional and phylogenetic diversity correlated at the global scale?

Functional and phylogenetic diversity are negatively correlated at the global scale

38



Positive correlation was shown before<sup>[19]</sup> Increasing with higher number of traits

Traits map differently on the phylogeny at smaller spatial extent<sup>[6, 7]</sup>





MARTIN-LUTHER-UNIVERSITÄT Georg J. A. Hähn HALLE-WITTENBERG



Discussion – To what extent is the geographical distribution of functional and phylogenetic diversity explained by present and past climate conditions?

Distribution pattern of functional diversity depends on current climatic conditions.





Edinburgh, Scotland  $18^{\text{th}} - 21^{\text{st}}$  December 2022



MARTIN-LUTHER-UNIVERSITÄT Georg J. A. Hähn



Discussion – To what extent is the geographical distribution of functional and phylogenetic diversity explained by present and past climate conditions?

40

Distribution pattern of functional diversity depends on current climatic conditions.



Functional diversity can be linked to recent climate conditions on smaller spatial extent <sup>[20]</sup>

But local communities depend on local factors such as land-use<sup>[21]</sup> or soil properties<sup>[22]</sup>





MARTIN-LUTHER-UNIVERSITÄT Georg J. A. Hähn HALLE-WITTENBERG

BES 2022 Annual Meeting

Discussion – To what extent is the geographical distribution of functional and phylogenetic diversity explained by present and past climate conditions?

Distribution pattern of functional diversity depends on current climatic conditions.



Functional diversity can be linked to recent climate conditions on smaller spatial extent  $^{\mbox{[20]}}$ 

But local communities depend on local factors such as land-use<sup>[21]</sup> or soil properties<sup>[22]</sup>



Spatial pattern of phylogenetic diversity depends on past climatic events,  $\rightarrow$  positive correlation with climatic variability after LGM in the GAM

Phylogenetic turnovers in regions with high climatic changes after the LGM<sup>[23]</sup>





Outlook – Why do we care?



Edinburgh, Scotland

18<sup>th</sup> – 21<sup>st</sup> December 2022



The results can be used to identify phylogenetic and functional diverse hotspots

But more information is needed in some regions, e.g., global south.

Feel free to contact us!

Please help us spread the call!



BES 2022 Annual Meeting

# Many thanks to you,

Plot

the sPlot consortium and data contributors,

and our supporters and partners



Edinburgh, Scotland

18<sup>th</sup> – 21<sup>st</sup> December 2022

Georg Hähn



Francesco Sabatini E-Mail: <u>georg.haehn@idiv.de</u> Web: <u>georghaehn.netlify.app</u> Twitter: <u>twitter.com/GeorgHaehn</u> Mastodon:

) iDiv

GeorgHaehn@ecoevo.social



Gabriella Damasceno





Helge Bruelheide

E-Mail: <u>gabriella.damasceno@idiv.de</u> Web: <u>https://www.idiv.de/en/splot.html</u> Twitter: <u>twitter.com/sPlot\_iDiv</u>

鄂 sPlot