

# Root traits in the context of the TRY database

Jens Kattge, Sandra Diaz, Sandra Lavorel, Colin Prentice, Paul Leadley, Gerhard Bönisch, Arindam Banerjee, Farideh Fazayeli, Hanhuai Shan, Franziska Schrodte, Julia Joswig, Peter Reich, Christian Wirth and all members of the TRY initiative



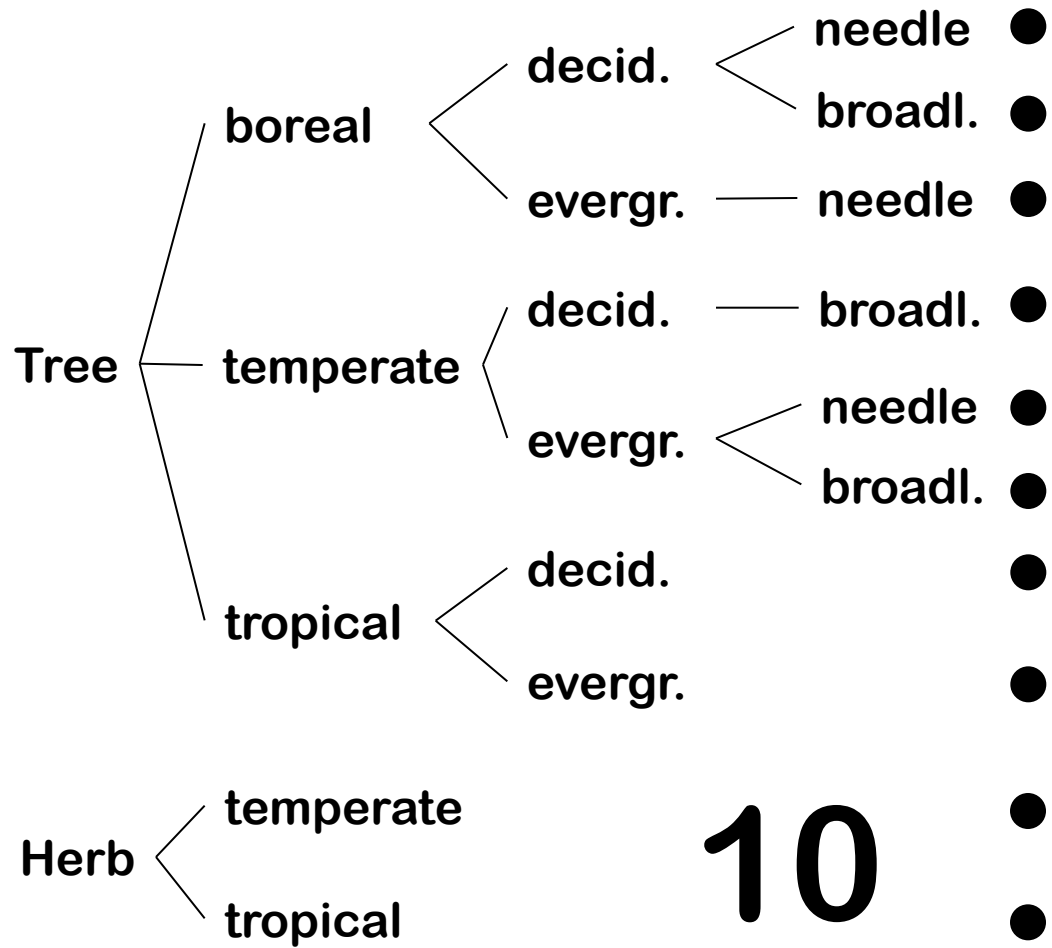
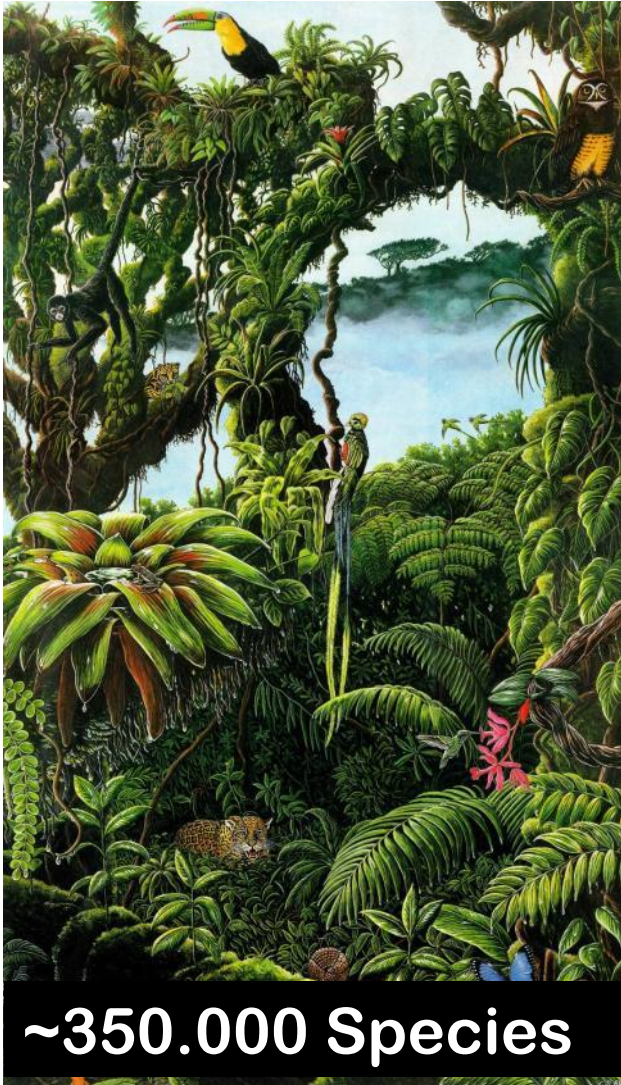
Max Planck Institute  
for Biogeochemistry



GLOBAL  
IGBP  
CHANGE  
International  
Geosphere-Biosphere  
Programme



# Diversity and DGVMs



# 2007



Fast Track Initiative on Refining Plant  
Functional Classifications



Sandra Díaz  
Sandra Lavorel  
Colin Prentice  
Paul Leadley

Christian Wirth  
Gerhard Bönisch  
Jens Kattge



**TRY**

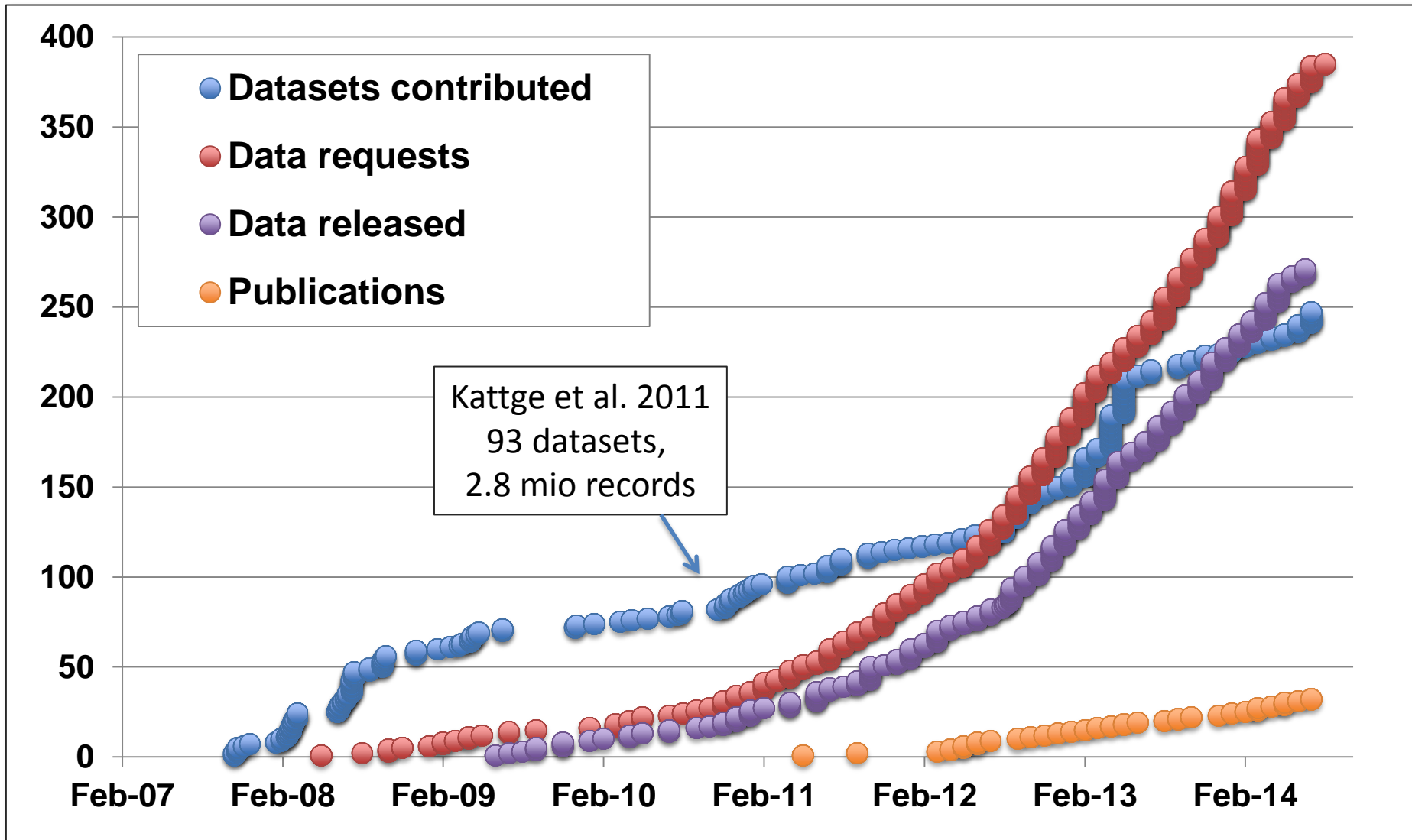
**Goal:** A global database of plant traits to make the data available for  
trait-based approaches in ecology and vegetation modeling

# Incentive driven data sharing

- **Data remain intellectual property of contributors**
  - Restricted access
  - Open access (new: autumn 2014)
- **Incentives for data sharing**
  - Use of (restricted) data from the joint database
  - Potential collaboration in publications
  - References being cited

**TRY**

# TRY has gained momentum

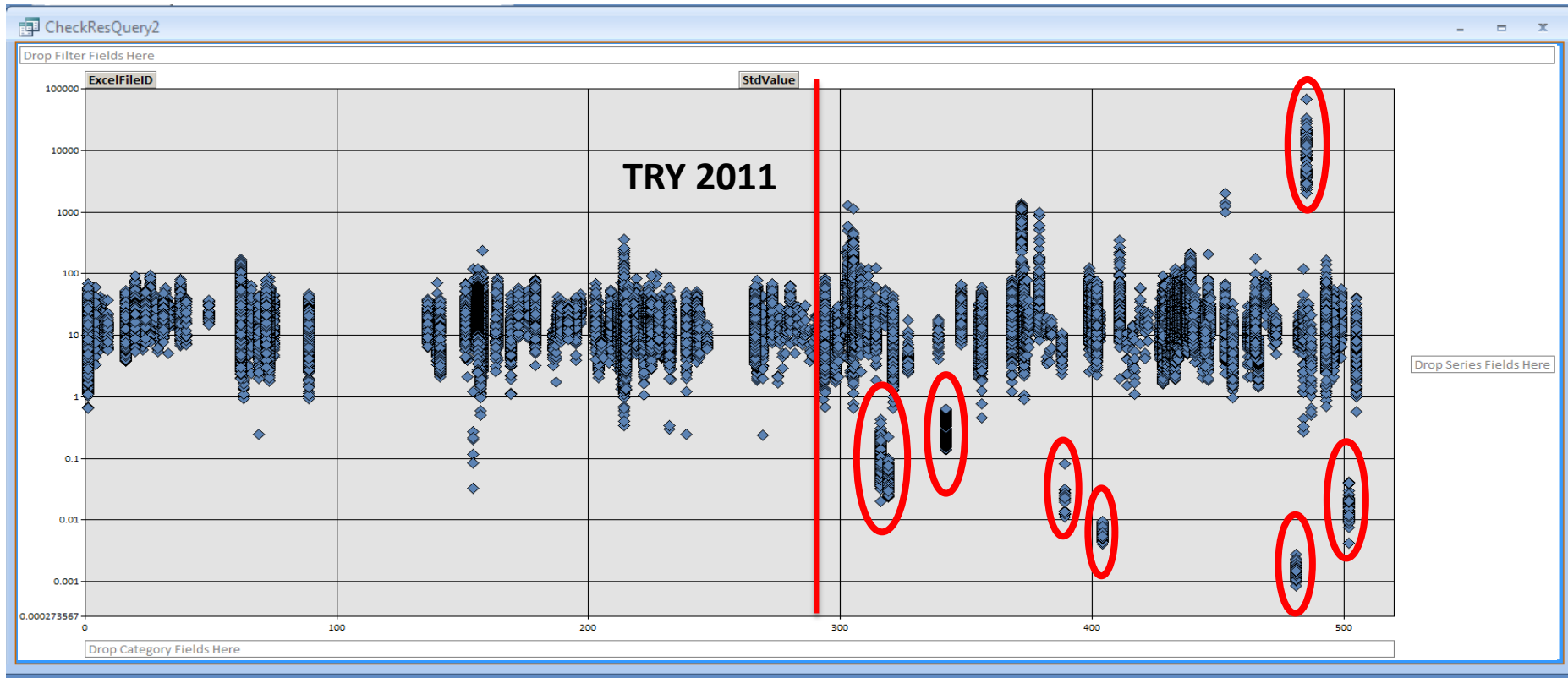


# Data processing & quality control

1. Data acquisition
2. Data import
3. Matching of concepts
  - Traits
  - Auxiliary data
  - Taxonomy
4. Standardization of units
5. Correction of errors (contributor's feedback)
6. Complementmentation of trait data
7. Identification of duplicates and outliers

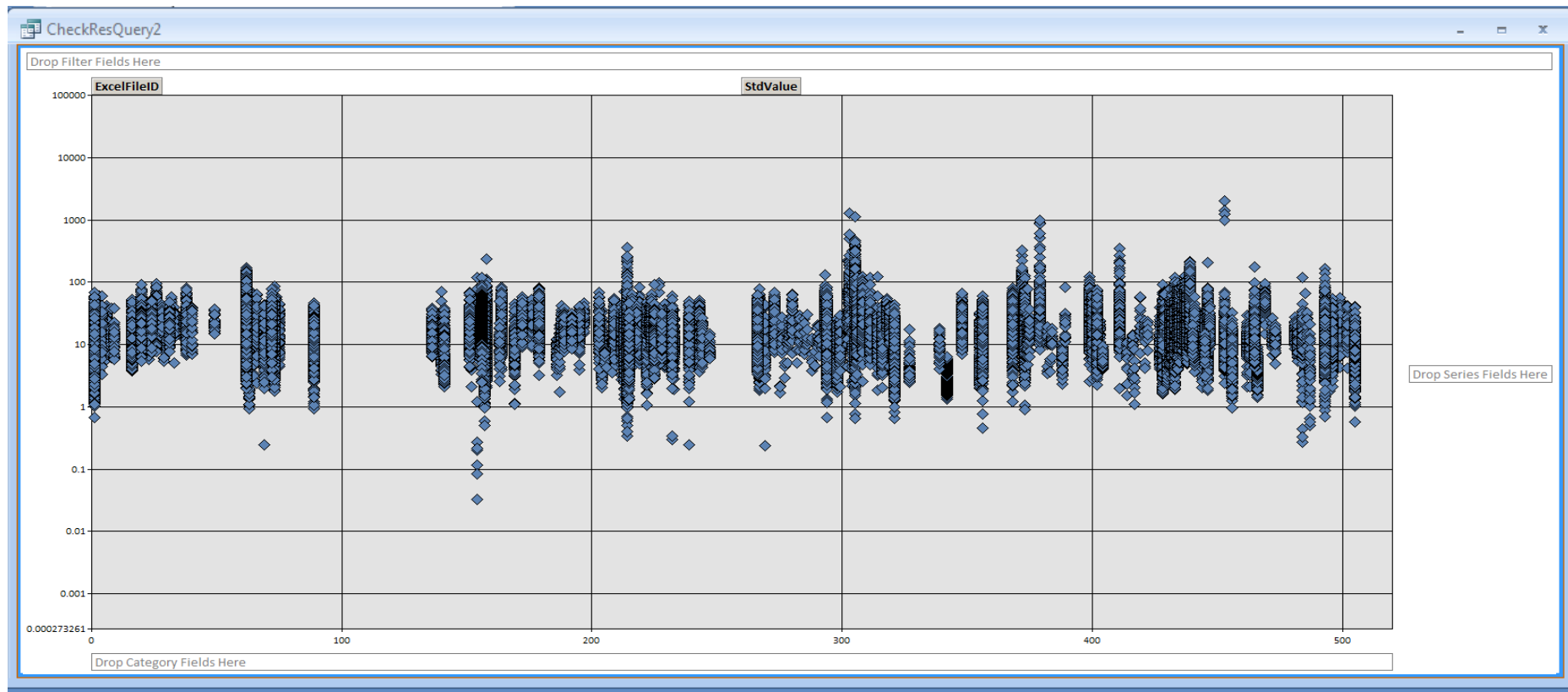
**TRY**

# Something's wrong here



e.g. specific leaf area (~90 new files)

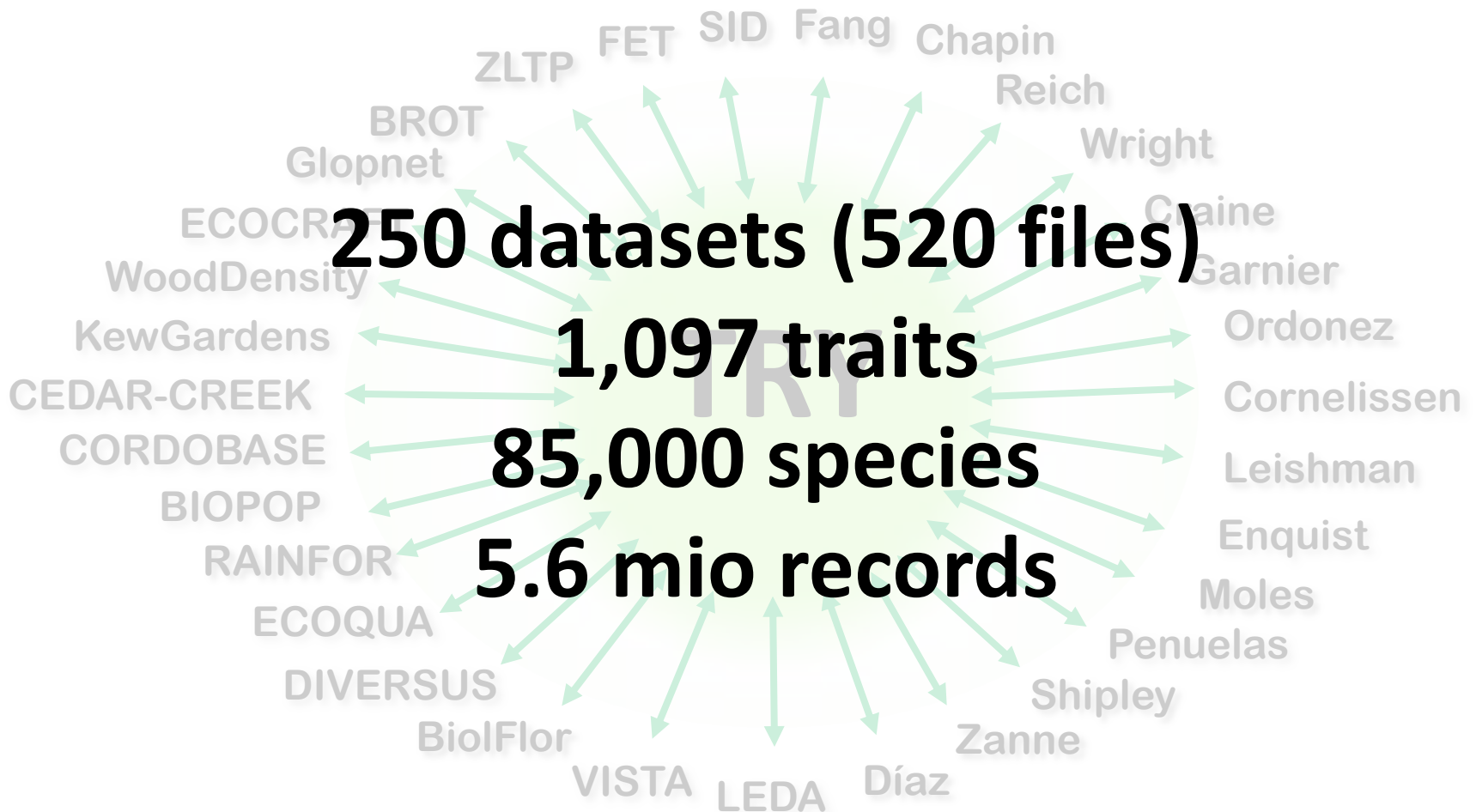
# ... after the correction of unit mismatches





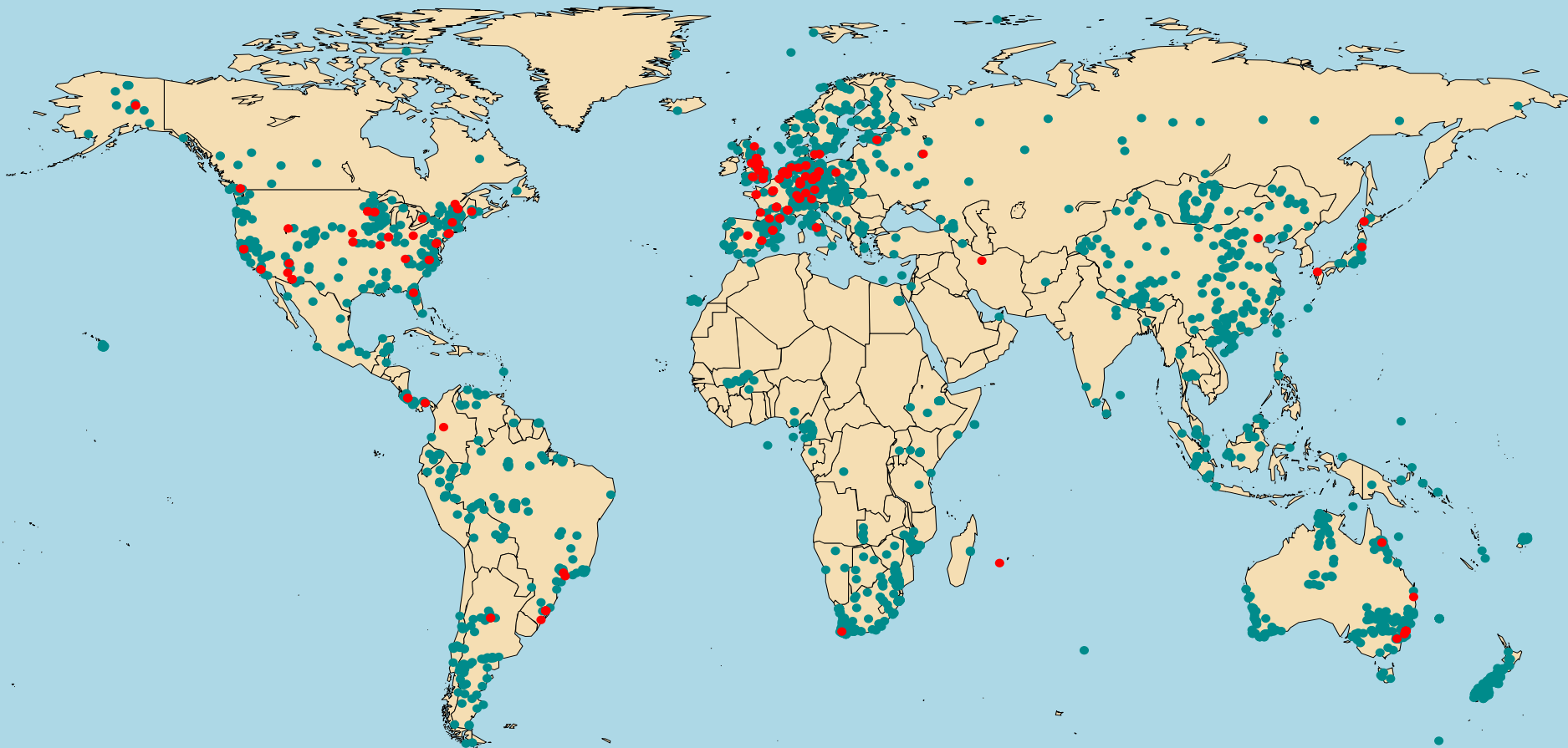
# TRY – a global database of plant traits

Second generation of data pooling

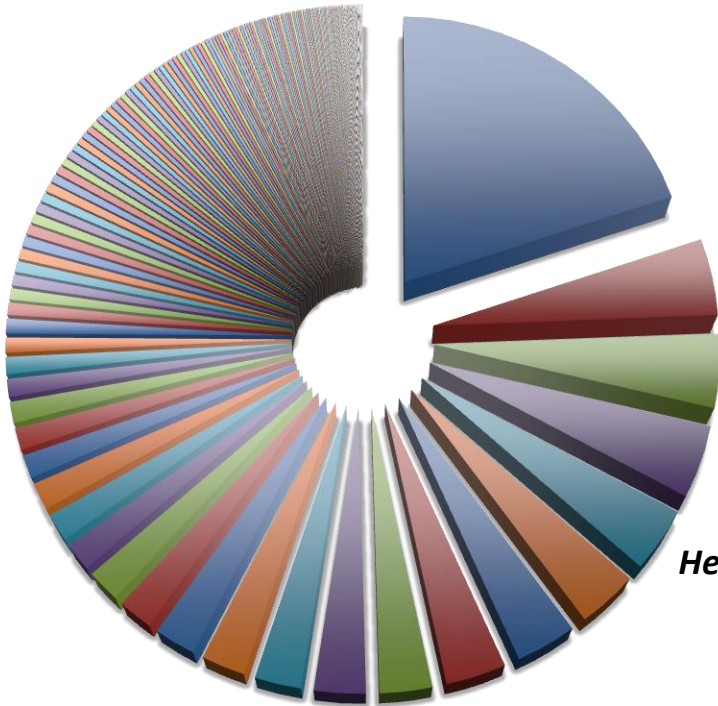


# TRY – a global research network

- 167 contributing partner institutes (372 scientists)
- 14361 measurement sites

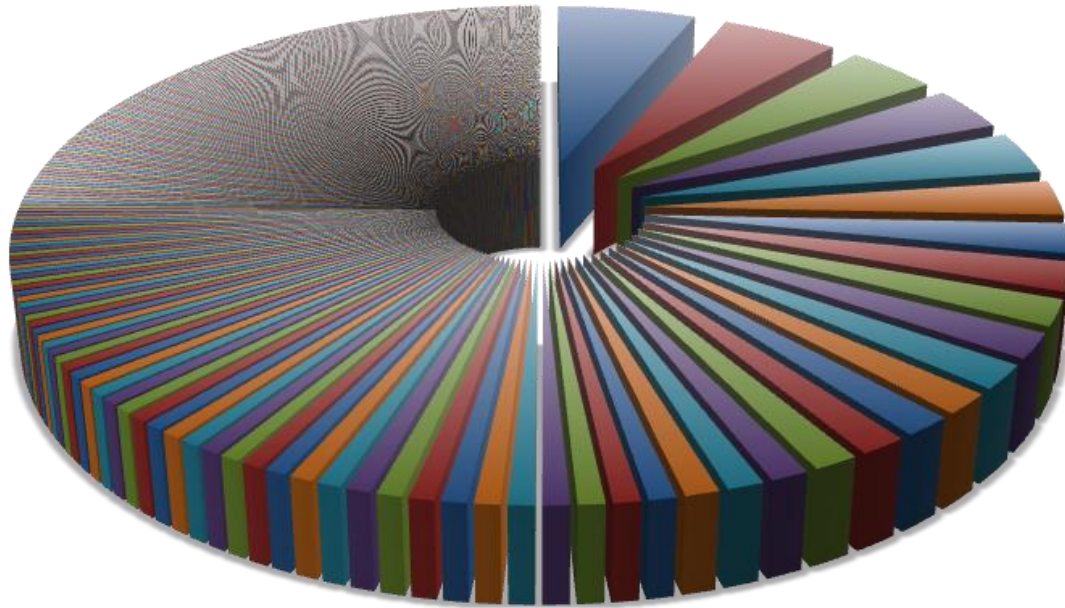


# Trait records by dataset



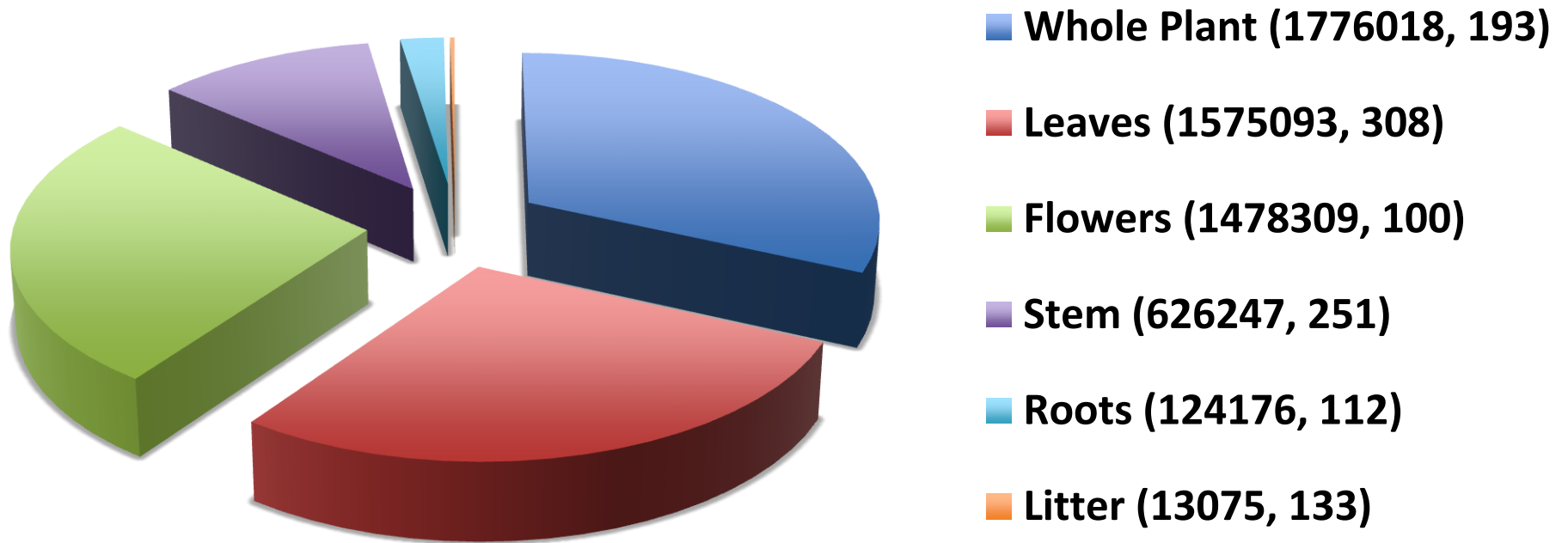
The LEDA Traitbase	1048680	Michael Kleyer
KEW Seed Information Database (SID)	238105	John Dickie
BioFlor Database	226840	Ingolf Kühn, Stefan Klotz
VegClass CBM Global Database	223588	Andy Gillison
<b><i>CLO-PLA - a database of clonal growth</i></b>	<b>196330</b>	<b><i>Jitka Klimesova</i></b>
PLANTSdata	178168	Walton Green
<b><i>Ecological Flora of the British Isles</i></b>	<b>177724</b>	<b><i>Alastair Fitter</i></b>
<b><i>The Functional Ecology of Trees (FET)</i></b>	<b>175329</b>	<b><i>C. Wirth, J. Kattge</i></b>
<b><i>Bridge database</i></b>	<b>152737</b>	<b><i>Christopher Baraloto</i></b>
<b><i>DBH and life form of Amazonian flora</i></b>	<b>149718</b>	<b><i>Erika Berenguer</i></b>
BIOPOP: Traits for Nature Conservation	136053	Peter Poschlod
<b><i>Chinese transect trait database</i></b>	<b>126805</b>	<b><i>Sandy Harrison</i></b>
<b><i>Herb water relations on moisture gradients</i></b>	<b>121918</b>	<b><i>Serge Sheremetev</i></b>
Panama Plant Traits Database	101487	S. Joseph Wright
<b><i>The xylem/phloem database</i></b>	<b>98726</b>	<b><i>Fritz Schweingruber</i></b>
Categorical Plant Traits Database	85027	Ian Wright
<b><i>Global woodiness database</i></b>	<b>84212</b>	<b><i>Amy Zanne</i></b>
<b><i>Plant Volatiles Database</i></b>	<b>84170</b>	<b><i>Almut Arneth</i></b>
Leaf structure and chemistry	71098	Bill Shipley
<b><i>D3: The Dispersal and Diaspore Database</i></b>	<b>67487</b>	<b><i>Oliver Tackenberg</i></b>
Global 15N Database	65051	Joseph Craine
GLOPNET - Global Plant Trait Network	55412	Ian Wright, Peter Reich
Reich-Oleksyn Global Leaf N, P Database	53724	Peter Reich
<b><i>Global Leaf Element Composition Database</i></b>	<b>46802</b>	<b><i>Steven Jansen</i></b>
KEW African Plant Traits Database	46253	Don Kirkup
Leaf Physiology Database	38804	Jens Kattge, C. Wirth

# Trait records by trait

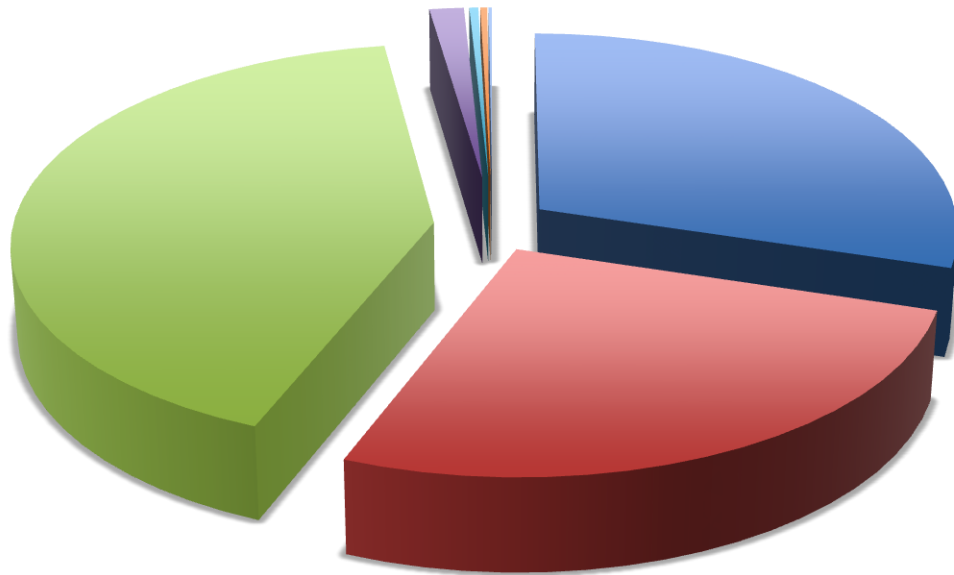


Dispersal syndrome	351826
Plant growth form	297906
<b>Plant height</b>	<b>229969</b>
<b>Seed dry mass</b>	<b>200770</b>
Dispersal unit type	195107
Dispersal unit floating capacity	159702
<b>Specific leaf area, SLA</b>	<b>137689</b>
<b>Leaf area</b>	<b>132172</b>
<b>Stem diameter</b>	<b>117567</b>
<b>Leaf dry matter content, LDMC</b>	<b>113031</b>
Plant woodiness	109148
Leaf phenology type	93358
Seedbank type	89951
<b>Leaf dry mass</b>	<b>87410</b>
Plant life form (Raunkiaer)	82710
<b>Budbank height distribution</b>	<b>70444</b>
<b>Leaf N content per leaf dry mass</b>	<b>67002</b>
<b>Seed germination efficiency</b>	<b>63202</b>
<b>Seed (seedbank) longevity</b>	<b>50959</b>
Plant reproductive phenology	50492

# Trait records by plant parts



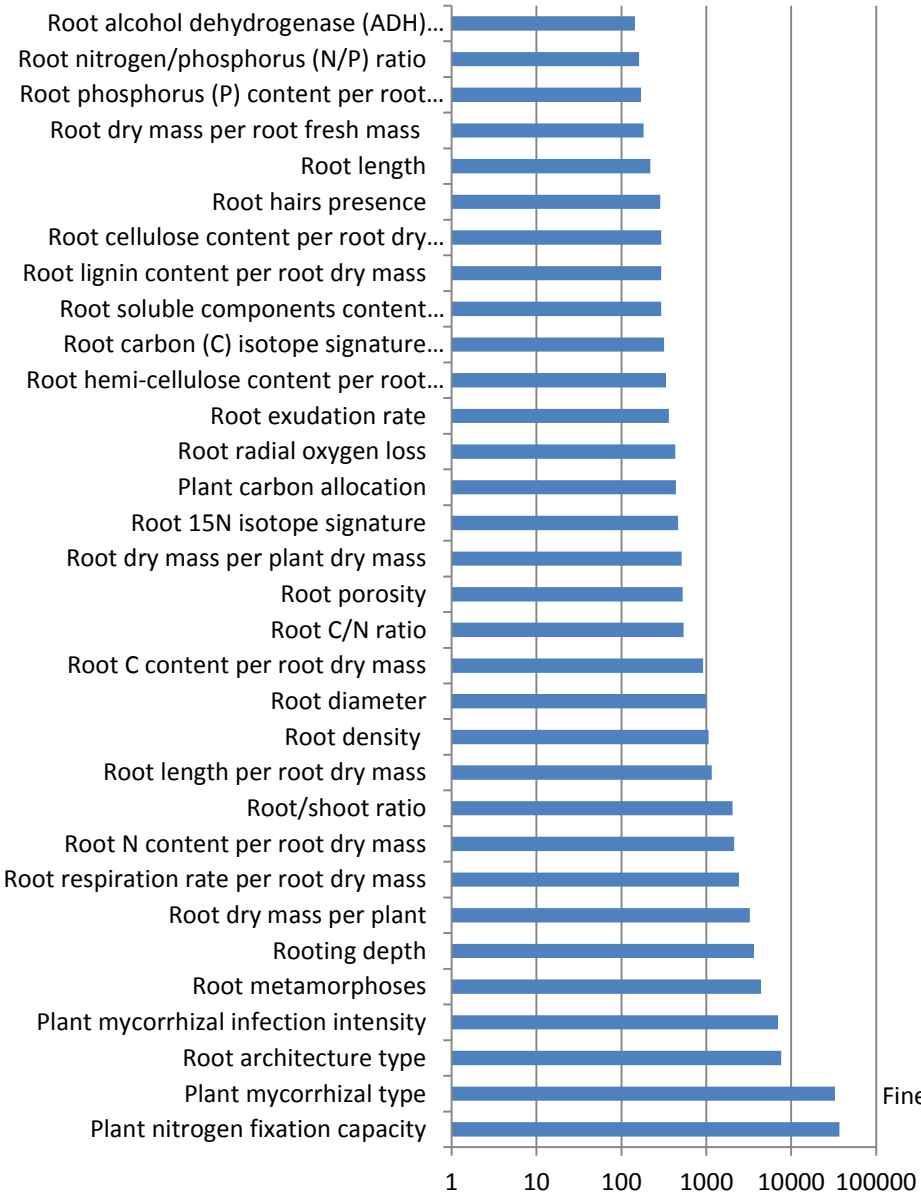
# Root traits in the context of TRY



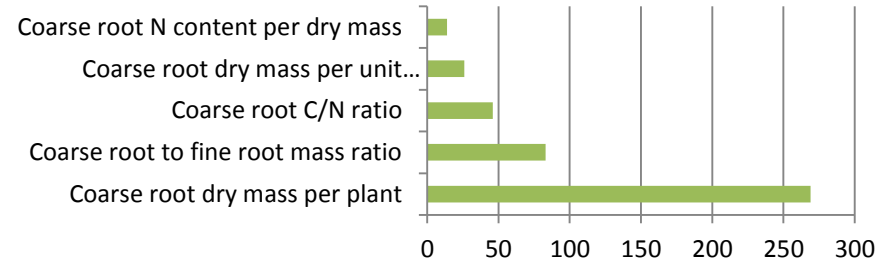
- Nitrogen fixation capacity (37081, 1)
- Mycorrhizal type (32797, 1)
- Roots in general (51959, 69)
- Fine roots (1859, 16)
- Coarse roots (480, 27)
- Coarse root CWD (371, 18)
- Fine root litter (187, 16)

# Root traits – a closer look

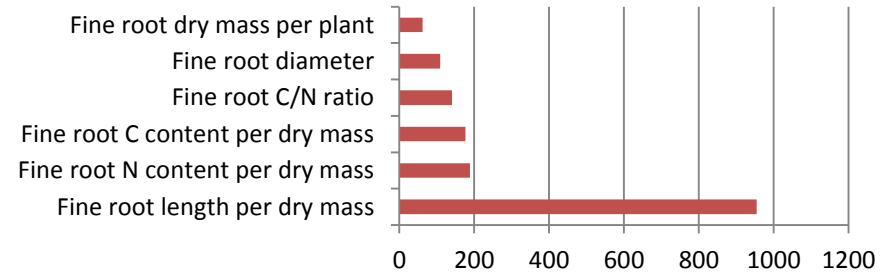
## Root traits (69)



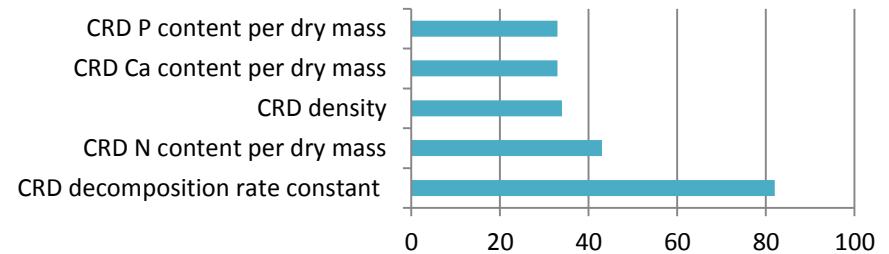
## Coarse roots (27)



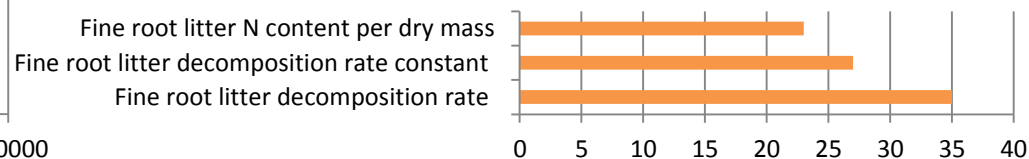
## Fine roots (16)



## Coarse root debris (18)



## Fine root litter (16)





# Gap-Filling of the TRY trait matrix

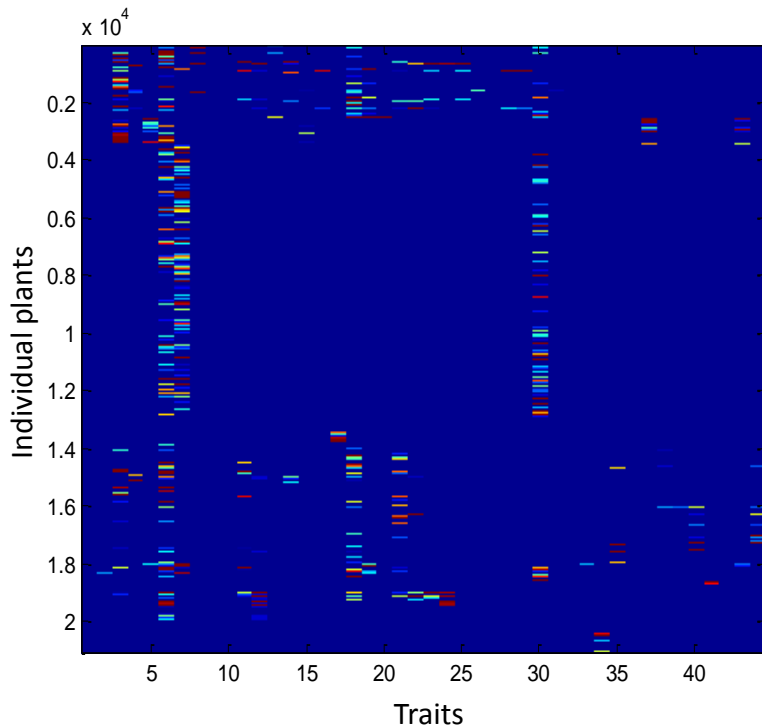


*a cooperation of the MPI-BGC and the University of Minnesota*

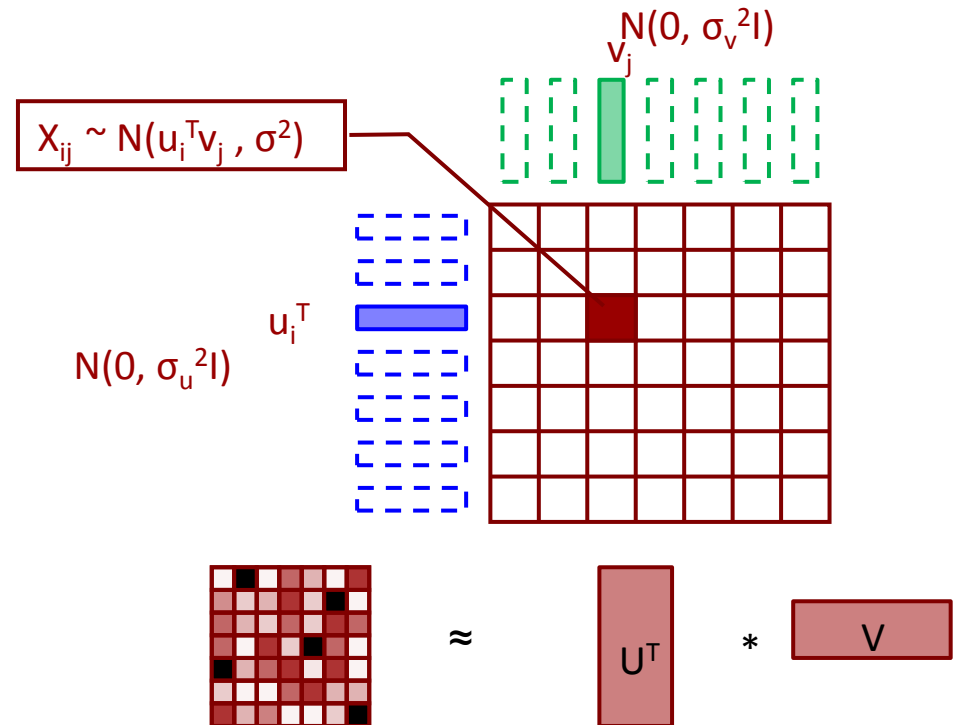
TRY = sparse matrix

5.6 million records on 1097 traits ×

2.2 million individuals (coverage 0.25 %)

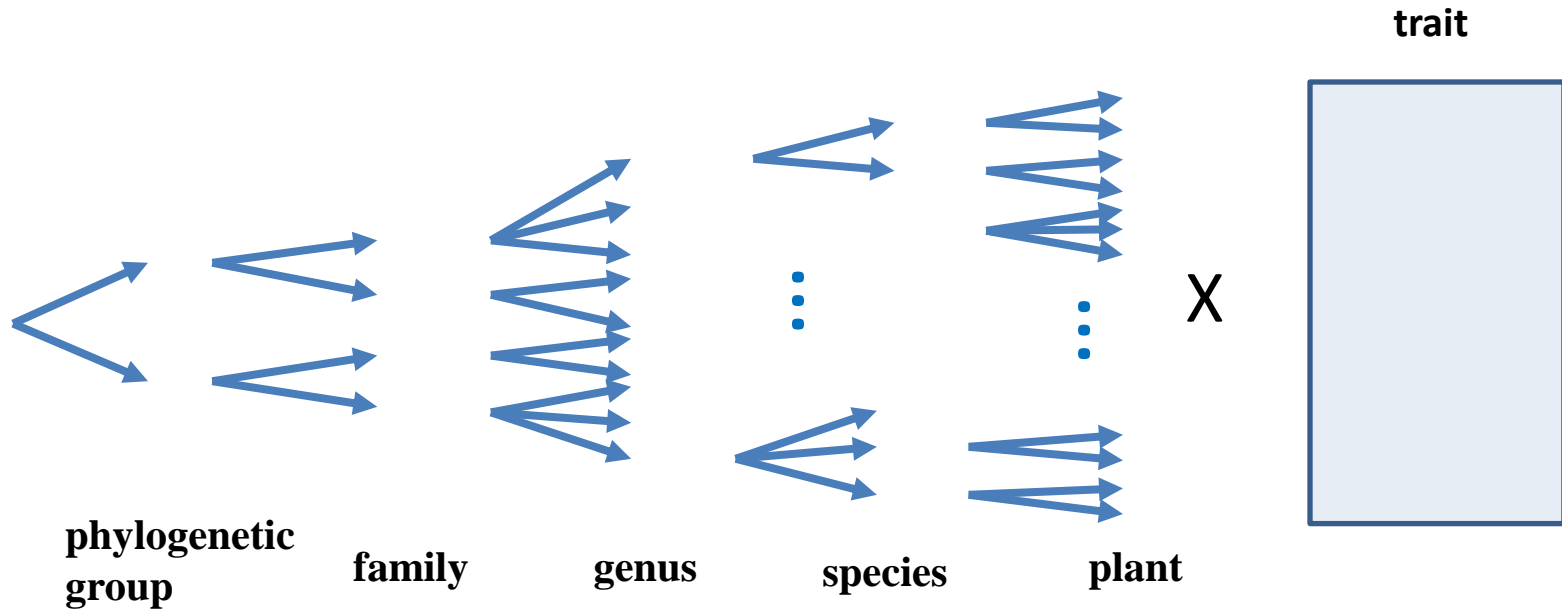


Probabilistic Matrix Factorization  
(PMF)





# Taxonomic hierarchy of the plant kingdom



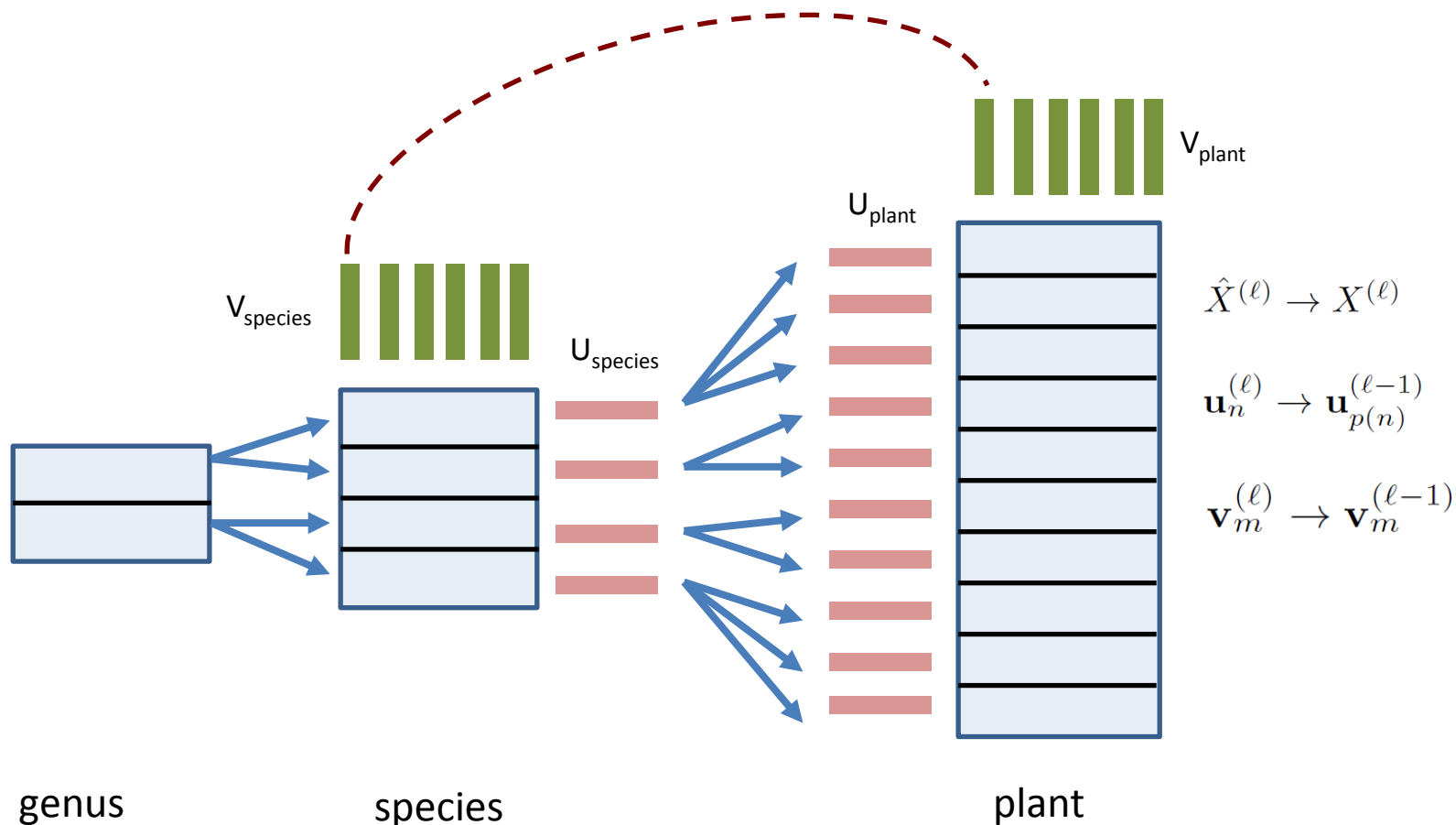
e.g. Scots pine

Gymnosperms      Pinaceae      Pinus      Pinus sylvestris



Plant height  
 Rooting depth  
 Leaf area  
 Leaf Mass  
 LMA  
 LeafN content  
 Photosynthesis rates  
 etc..

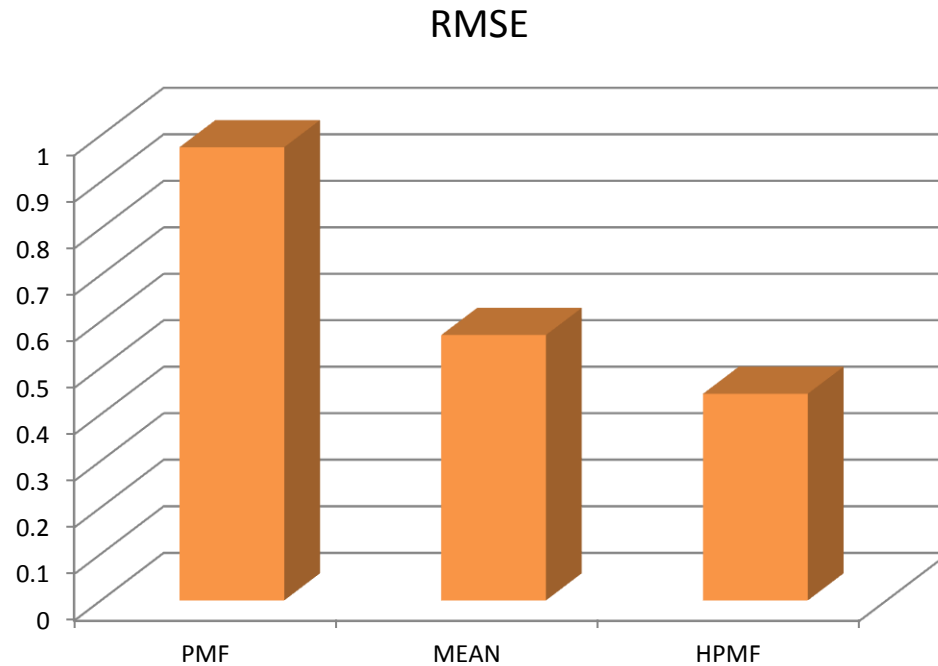
# Hierarchical Probabilistic Matrix Factorization



$$E = \sum_{\ell=1}^L \left\{ \sum_{nm} \delta_{nm}^{(\ell)} \left\| x_{nm}^{(\ell)} - \langle \mathbf{u}_n^{(\ell)}, \mathbf{v}_m^{(\ell)} \rangle \right\|_2^2 + \lambda_u \sum_n \left\| \mathbf{u}_n^{(\ell)} - \mathbf{u}_{p(n)}^{(\ell-1)} \right\|_2^2 + \lambda_v \sum_m \left\| \mathbf{v}_m^{(\ell)} - \mathbf{v}_m^{(\ell-1)} \right\|_2^2 \right\}$$

# HPMF compared to other algorithms

- 8 phylogenetic groups, 450 families, 7160 genera, 45824 species, 273777 plants, 443423 measurements
- 18 traits

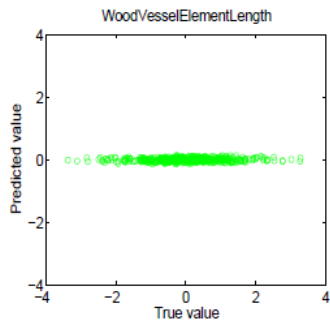


•HPMF outperforms PMF and MEAN

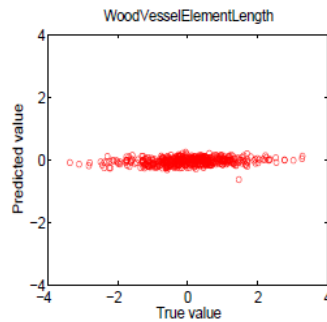
# (true value, predicted value) scatter plot

## Wood Vessel Element Length

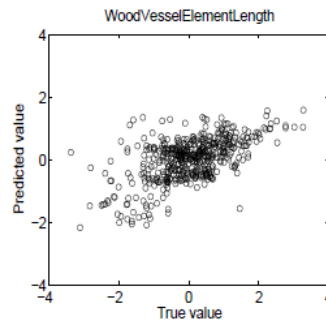
no phylo. info.



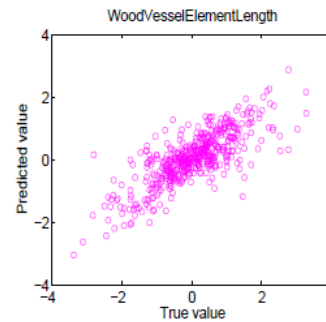
phylo.-group



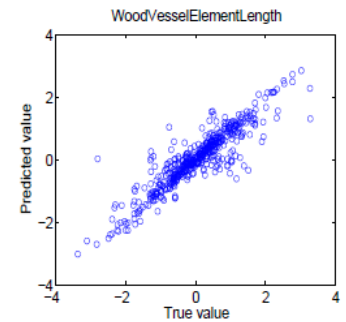
family



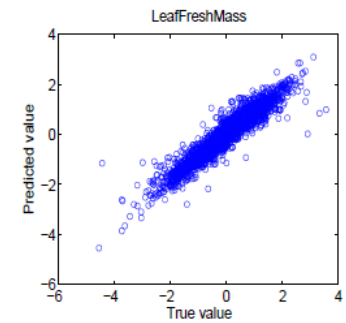
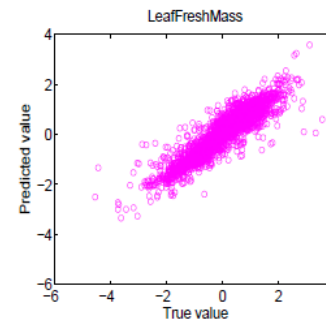
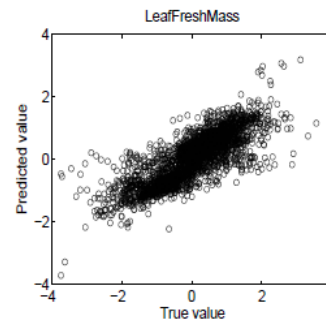
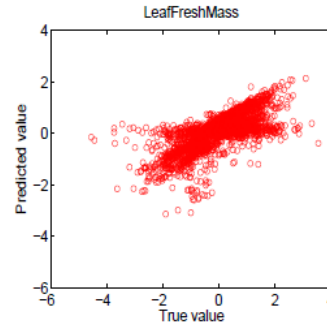
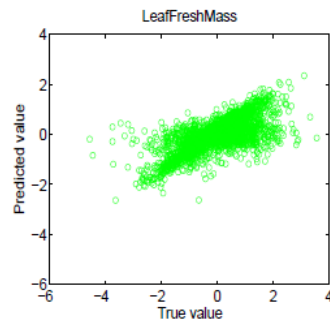
genus



species



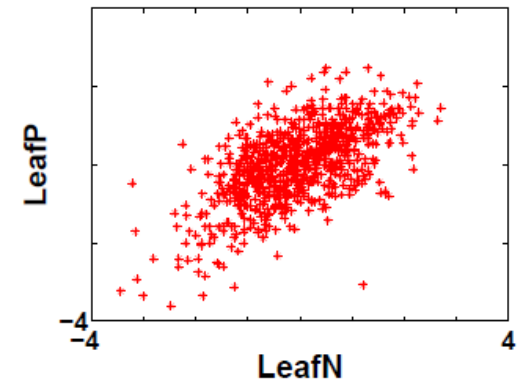
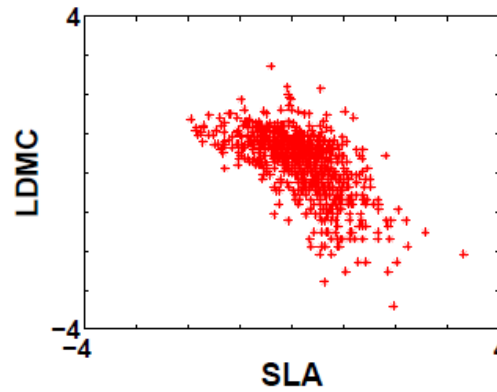
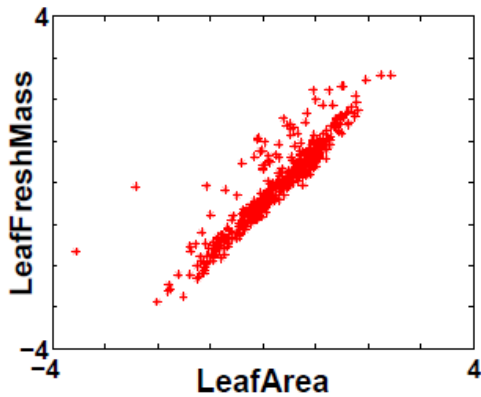
## Leaf Fresh Mass



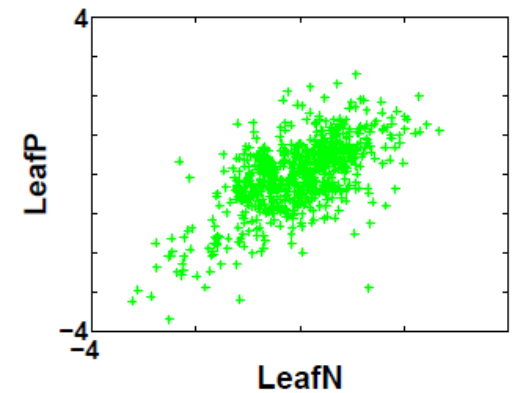
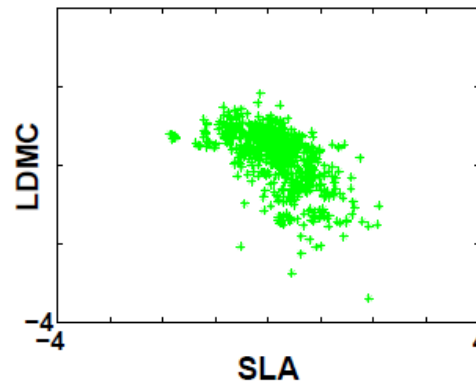
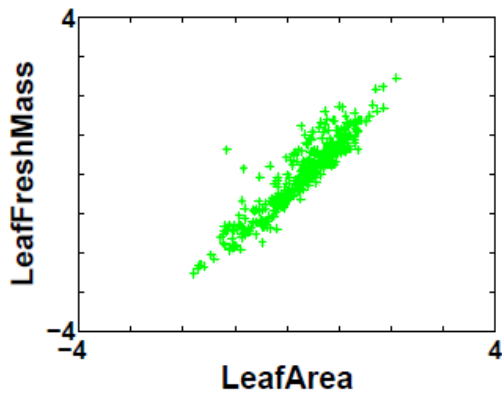
With increasing amount of taxonomic information, the plot becomes closer to a 1:1 line  
Deviations of true and predicted values indicate outliers

# Correlation between traits

true

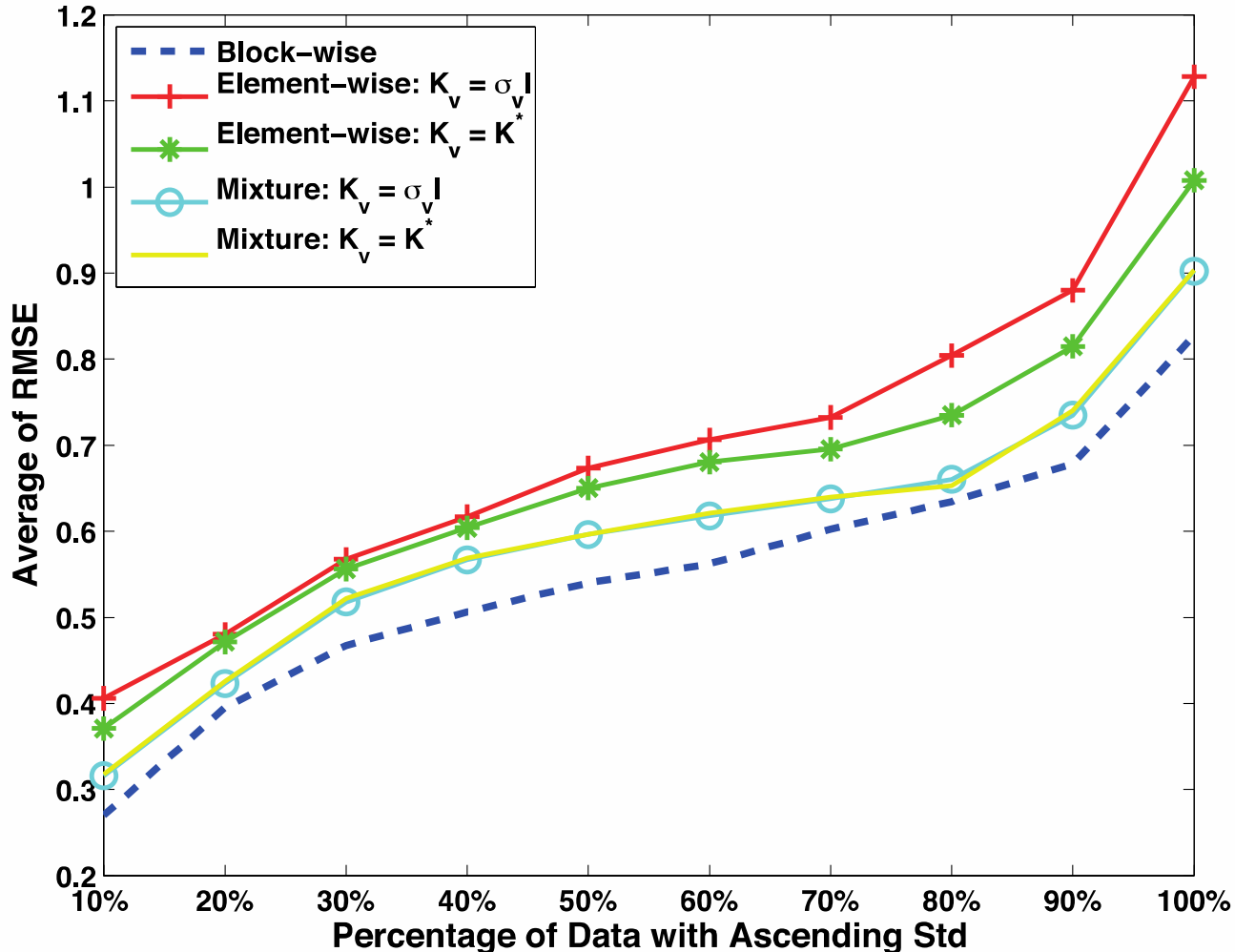


predicted



- The predicted trait correlation is close to the true plot

# Prediction error (RMSE) correlates with prediction uncertainty (Std)



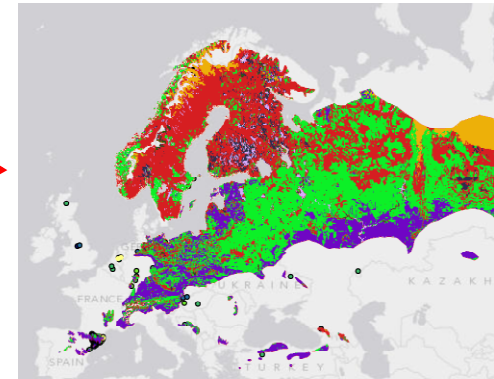
# Outlook: environment informed trait prediction for vegetation modeling

*a cooperation of the MPI-BGC, the University of Minnesota and the Oak Ridge National Laboratory*

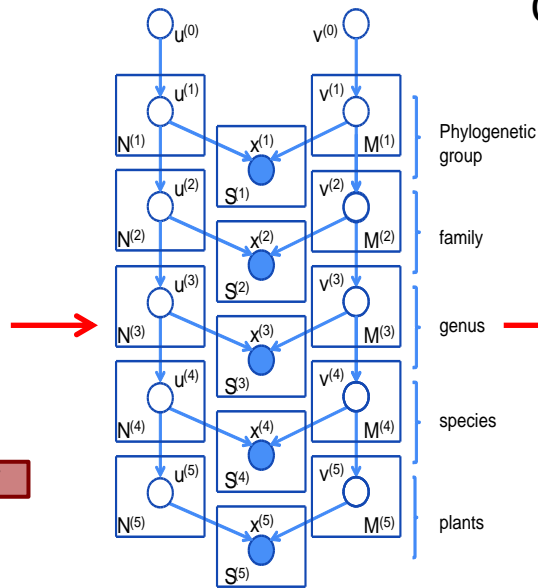
## Environment

Climate (e.g. WorldClim)  
Soil (e.g. HWSD)

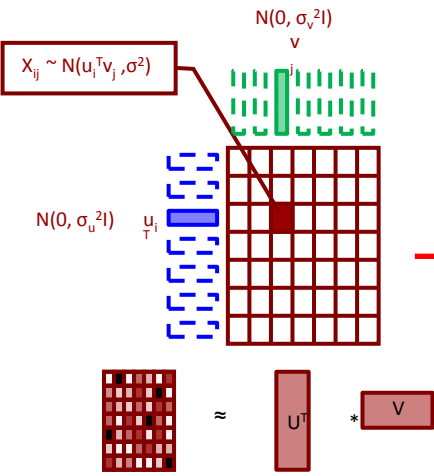
Regression  
against  
residuals  
or  
gap-filled  
data



Out of  
Sample  
Trait  
Prediction



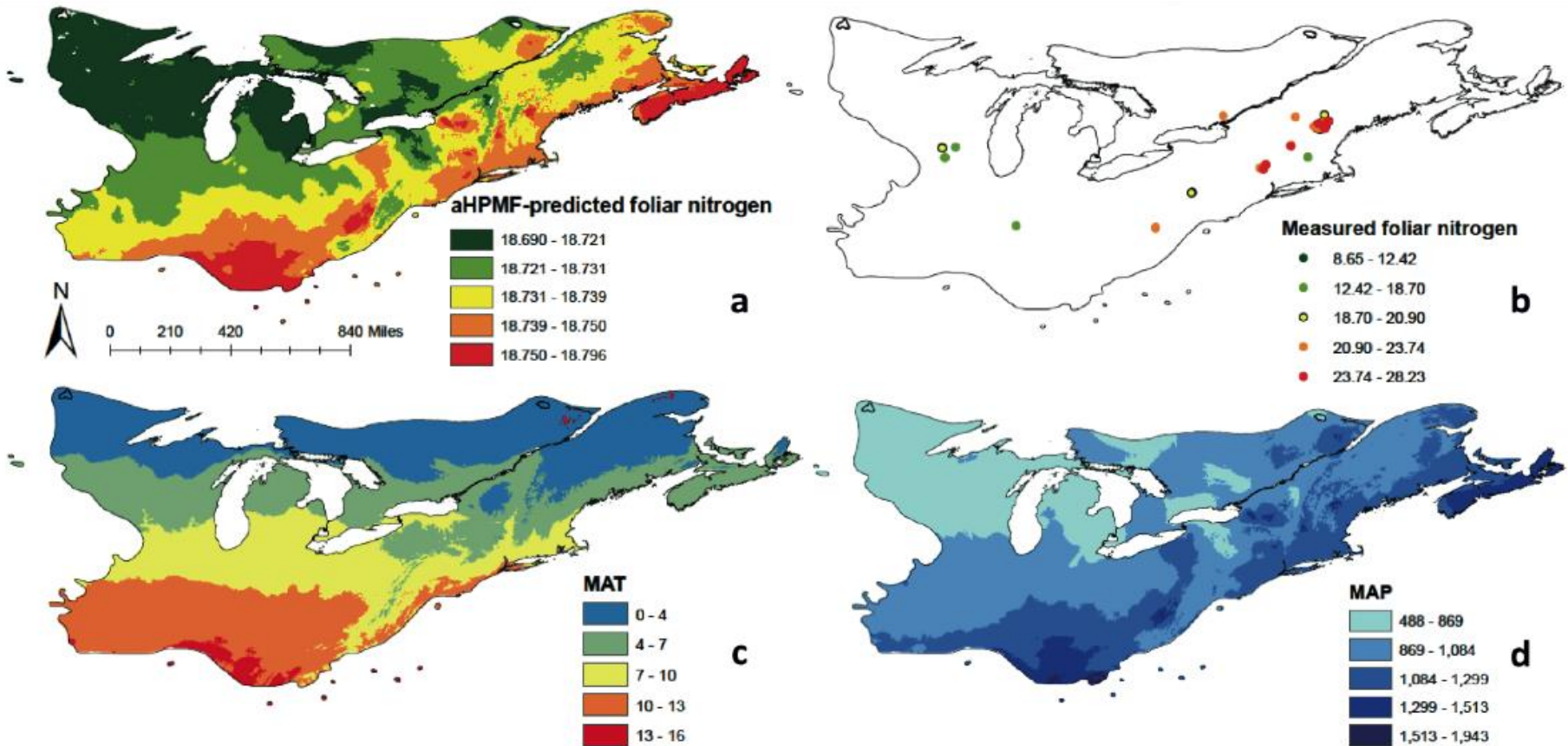
Hierarchical  
PMF



Probabilistic  
Matrix  
Factorization

# Out of Sample Prediction

e.g. Leaf N concentration for *Acer saccharum*





# Summary: Root Traits in TRY

1. **Data coverage in TRY doubled** from 2.8 million trait records in 2011 to 5.6 in 2014, **major part of the data will become publicly available** (autumn 2014)
2. All plant parts are almost equally well represented, **roots are underrepresented** and split to several traits, most with few entries each
3. The unprecedented coverage of the TRY database provides a **unique opportunity for gap-filling, quality assurance and trait prediction**

# TRY

# Plant Trait Database

Col. Vidalis, Cava  
61 traits  
Photo by A. Thuille

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## Quantifying and scaling global plant trait diversity

TRY is a network of vegetation scientists headed by [DIVERSITAS](#), [IGBP](#), the Max Planck Institute for Biogeochemistry and an international [Steering Committee](#).

### Main objectives

- Provide a global archive of plant traits
- Promote trait-based approaches in ecology and biodiversity science
- Support the design of a new generation of global vegetation models

### Current state of database and network

- 3 million trait records for about 69000 plant species
- 591 participants from 207 scientific institutes worldwide
- 256 scientific projects requesting plant trait data from TRY



### News

**Paper published** (2014-08-01)  
Walker AP et al.: The relationship of leaf photosynthetic traits – Vcmax and Jmax – to leaf nitrogen, leaf phosphorus, and specific leaf area: a meta-analysis and modeling study. *Ecology and Evolution* ([link](#))

**Paper published** (2014-06-23)  
Scheffer et al.: Why trees and shrubs but rarely trubs? *Trends in Ecology & Evolution*. ([link](#))

**Paper published** (2014-06-13)  
Werner et al.: A single evolutionary innovation drives the deep evolution of symbiotic N2-fixation in angiosperms. *Nature Communications*. ([link](#))

**Paper published** (2014-05-16)  
Vogel AT et al.: Which is a better predictor of plant traits: temperature or precipitation? *Journal of Vegetation Science*. ([link](#))

**Paper published** (2014-04-26)  
van Meirvenne et al.: Plant trait analysis delivers a comprehensive list of potential green roof species for the Mediterranean. *Ecological Engineering* ([link](#))

[News Archive](#)

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# www.try-db.org

