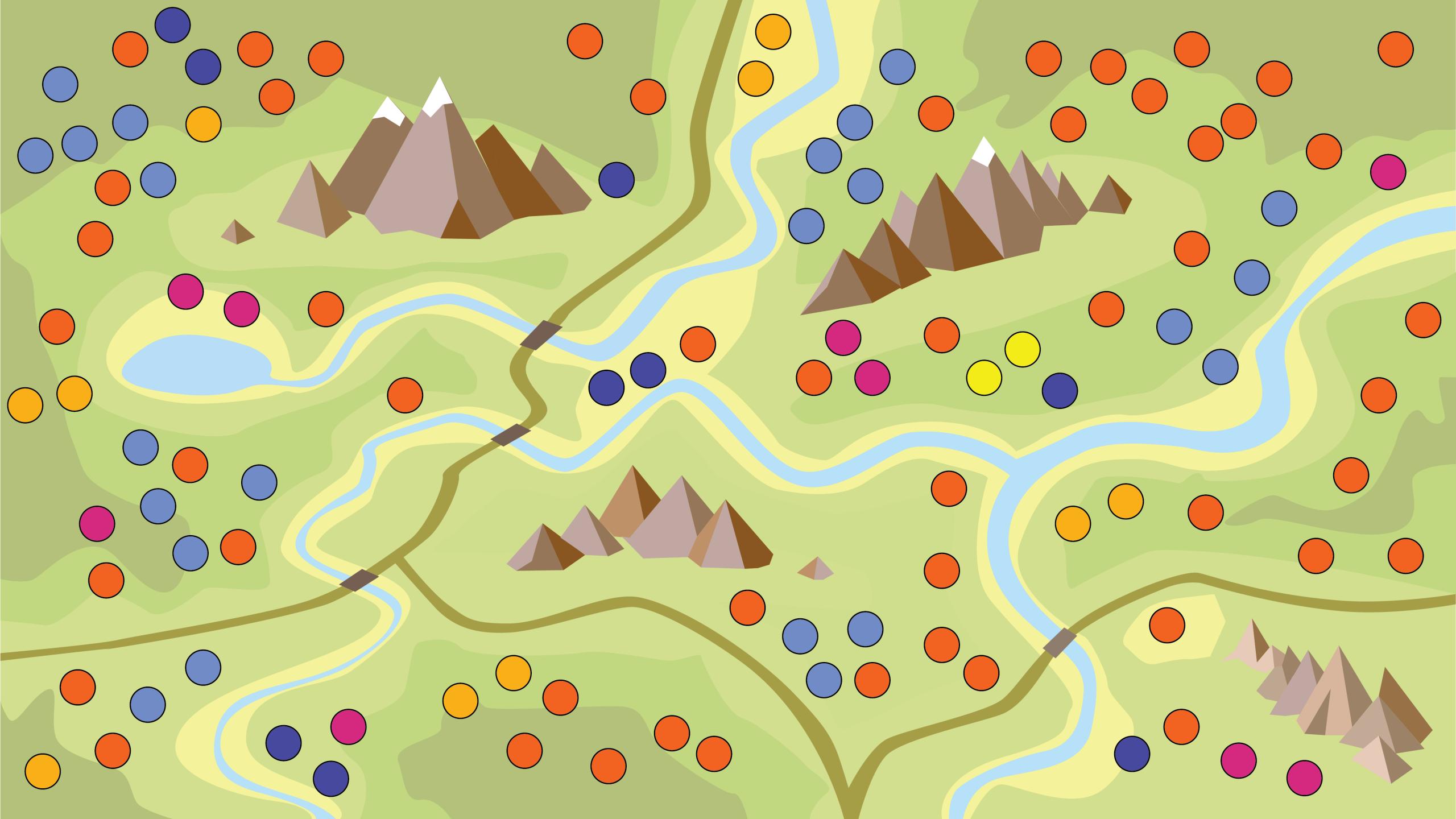
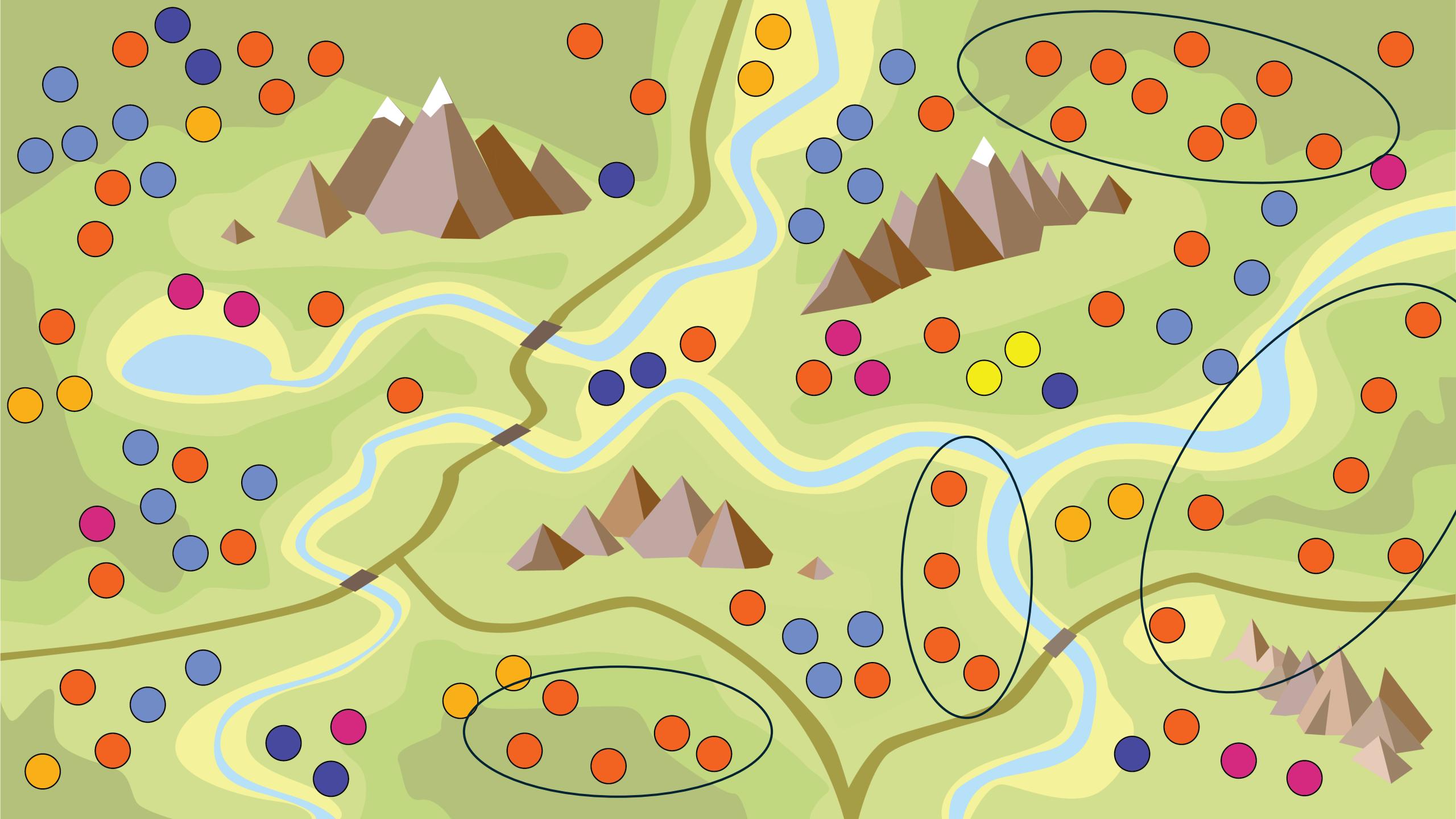


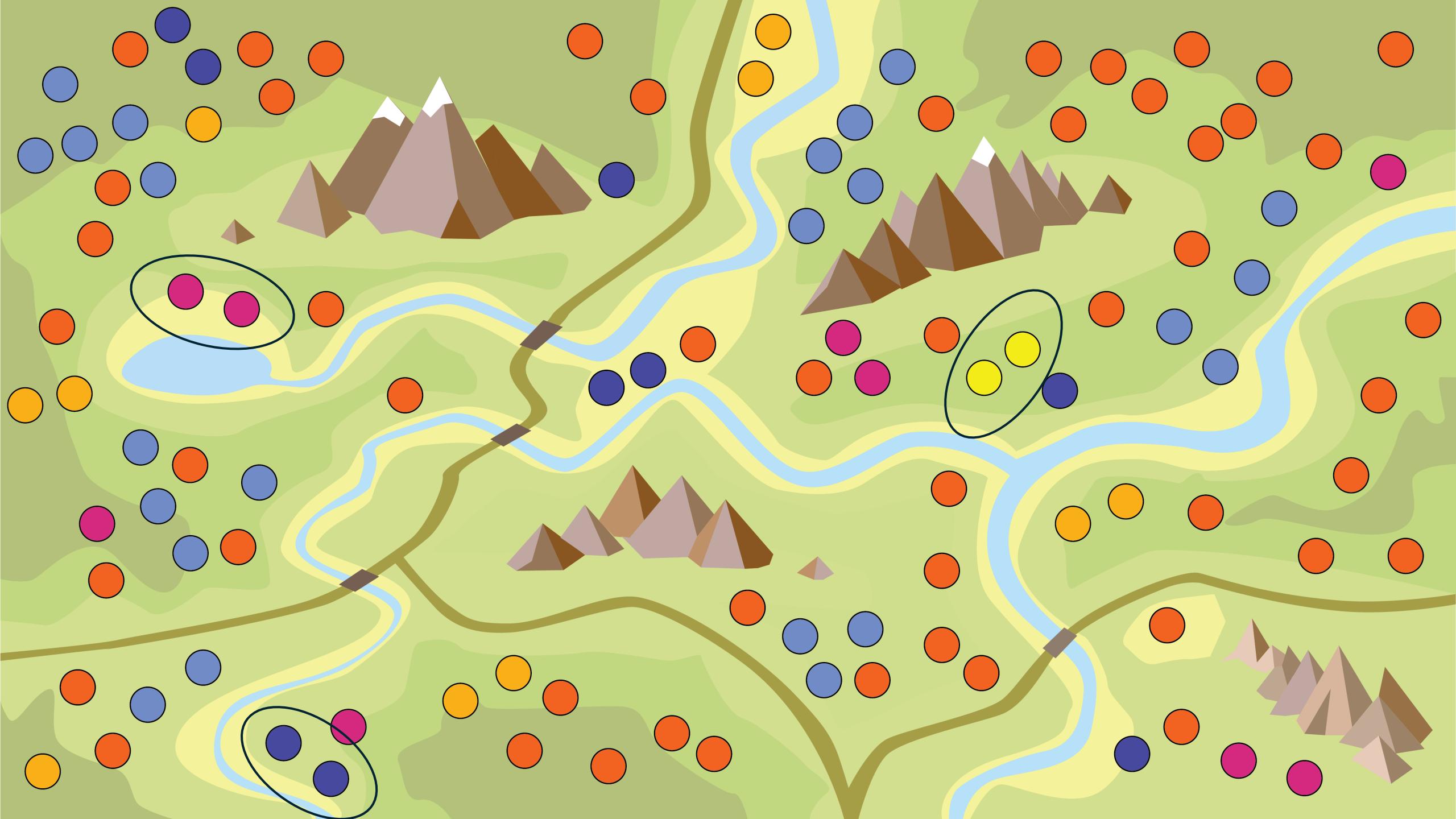
A Global Model of Discretized Rarity



Alivia G. Nytko, Brian C. O'Meara,
Joe K. Bailey









Geographic Range

Large

Small

Habitat Specificity

Wide

Narrow

Wide

Narrow

**Local Population
Size**

Large

Small

Geographic Range

Large

Small

Habitat Specificity

Wide

Narrow

Wide

Narrow

**Local Population
Size**

Large

Small

			Endemic

Geographic Range

Large

Small

Habitat Specificity

Wide

Narrow

Wide

Narrow

**Local Population
Size**

Large

Small

	Sparse		

Geographic Range	Large	Small		
Habitat Specificity	Wide	Narrow	Wide	Narrow
Local Population Size	Large			
	Small			

Functional Rarity: Trait Distinctiveness

Phylogenetic Rarity: Evolutionary Distinctiveness

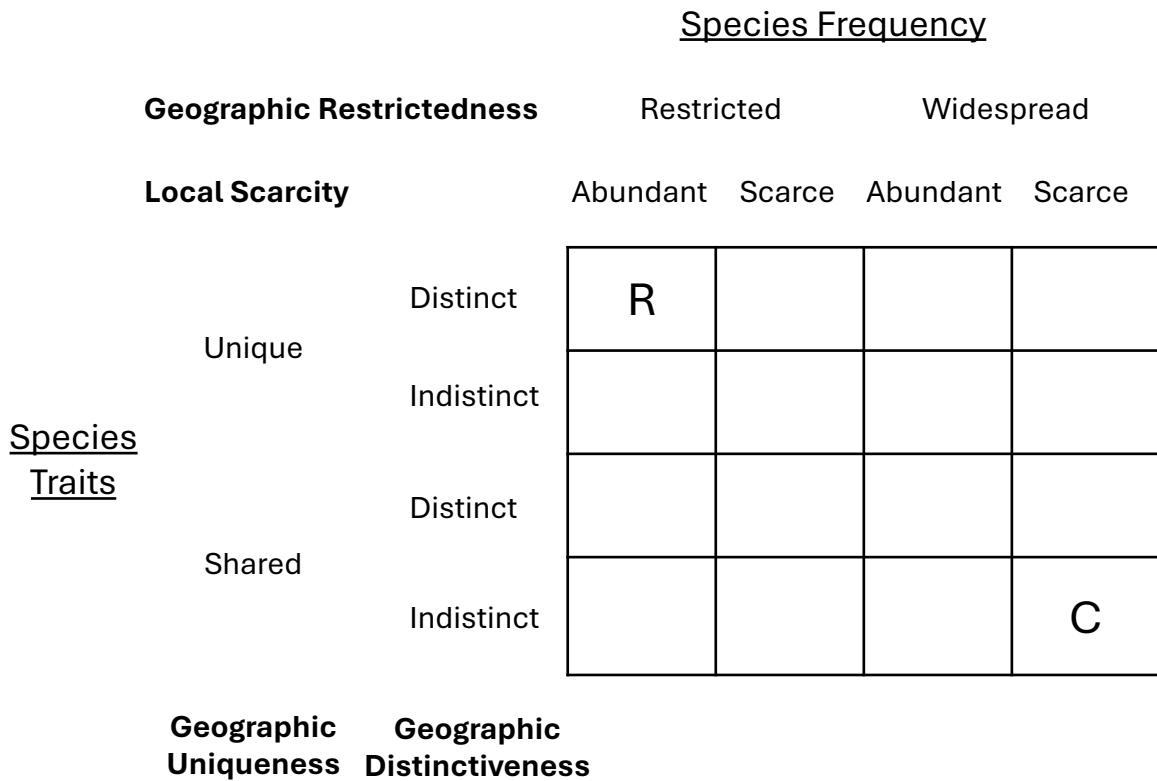
Geographic Range	Large		Small	
Habitat Specificity	Wide	Narrow	Wide	Narrow
Local Population Size	Large			
	Small			

Functional Rarity: Trait Distinctiveness

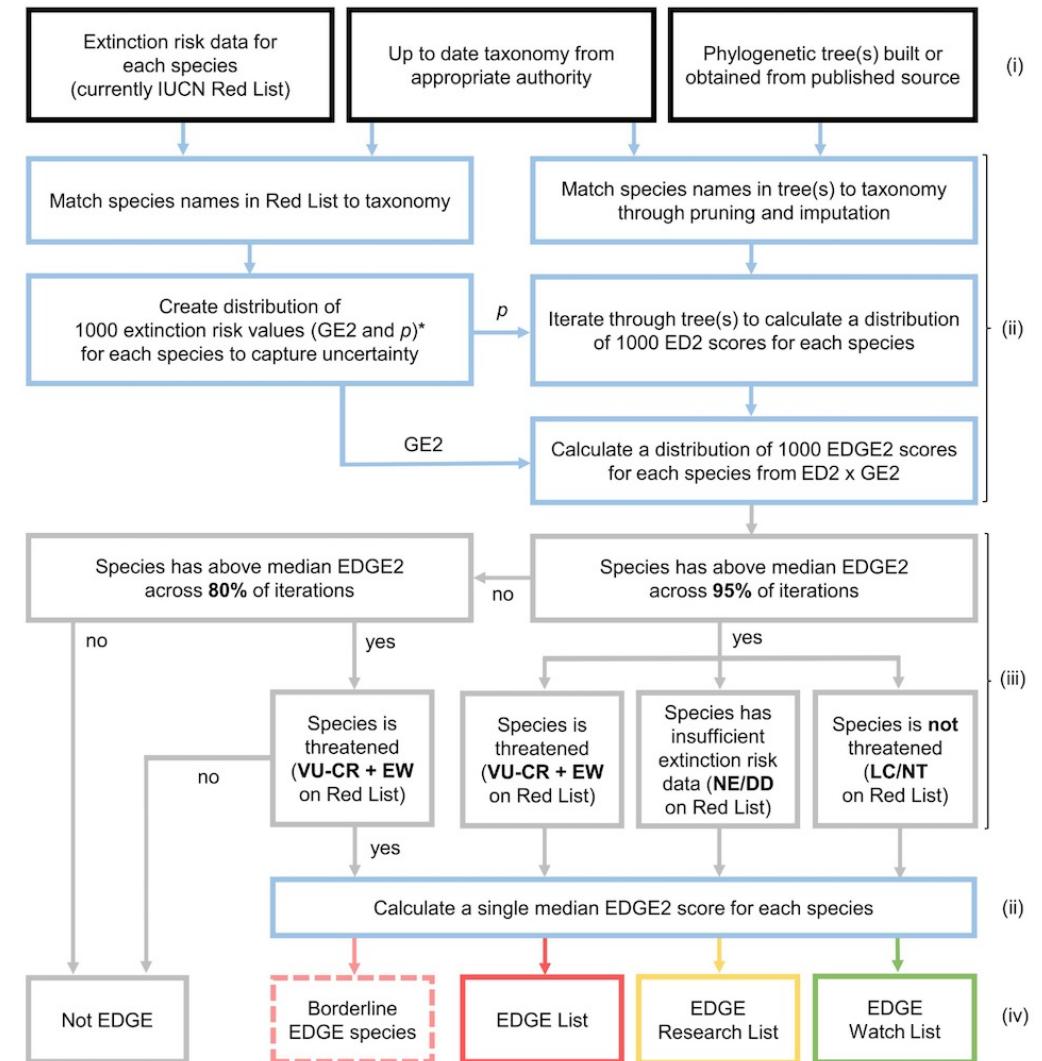
Phylogenetic Rarity: Evolutionary Distinctiveness

Geographic Range	Large		Small	
Habitat Specificity	Wide	Narrow	Wide	Narrow
Local Population Size	Large			
Small				

Functional Rarity Framework



EDGE Protocol



Functional Rarity Framework

EDGE Protocol

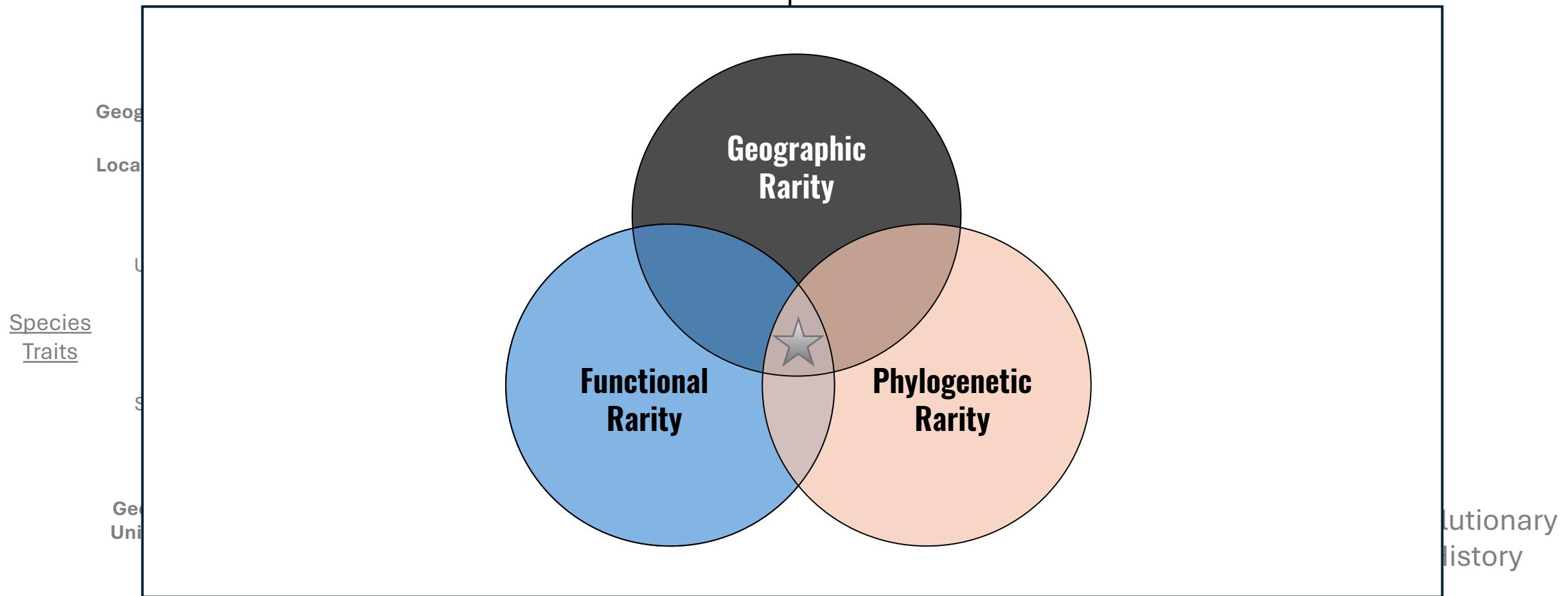


Table 1

Summary of the links between rarity, distinctiveness, originality, and diversity developed in the main text with diversity indices Q , ${}^{\alpha}K$ and ${}^{\alpha}K^*$.

	Quadratic diversity ^a	First parametric extension ^a	Second parametric extension ^a
Species-level			
FP-distinctiveness	$D_j = \sum_{i \neq j}^N \frac{p_i}{1 - p_j} d_{ij}$	${}^{\alpha}D_j = \frac{1 - \left(\sum_{i=1}^N p_i s_{ij} \right)^{\alpha-1}}{1 - p_j^{\alpha-1}}$	${}^{\alpha}D_j^* = \frac{\sum_{c=1}^{N-1} u_{c j} \left(1 - p_{c j}^{\alpha-1} \right)}{\left(1 - p_j^{\alpha-1} \right)}$
Abundance-based rarity	$\rho_j = (1 - p_j)$	${}^{\alpha}\rho_j = \frac{1 - p_j^{\alpha-1}}{\alpha - 1}$	${}^{\alpha}\rho_j = \frac{1 - p_j^{\alpha-1}}{\alpha - 1}$
Effective originality	$O_j = U_j \times \rho_j$	${}^{\alpha}O_j = {}^{\alpha}D_j \times {}^{\alpha}\rho_j$	${}^{\alpha}O_j^* = {}^{\alpha}D_j^* \times {}^{\alpha}\rho_j$
Community-level			
Species diversity ^b	$S = \sum_{j=1}^N p_j \rho_j$ $= 1 - \sum_{i=1}^N p_i^2$	${}^{\alpha}S = \sum_{j=1}^N p_j \left({}^{\alpha}\rho_j \right)$ $= \frac{1 - \sum_{j=1}^N p_j^{\alpha}}{\alpha - 1}$	${}^{\alpha}S = \sum_{j=1}^N p_j \left({}^{\alpha}\rho_j \right)$ $= \frac{1 - \sum_{j=1}^N p_j^{\alpha}}{\alpha - 1}$
FP-diversity	$Q = \sum_{j=1}^N p_j O_j$ $= \sum_{i=1}^N \sum_{j=1}^N p_i p_j d_{ij}$	${}^{\alpha}K = \sum_{j=1}^N p_j \left({}^{\alpha}O_j \right)$ $= \sum_{j=1}^N p_j \times \frac{1 - \left(\sum_{i=1}^N p_i s_{ij} \right)^{\alpha-1}}{\alpha - 1}$	${}^{\alpha}K^* = \sum_{j=1}^N p_j \left({}^{\alpha}O_j^* \right)$ $= \sum_{j=1}^N p_j \times \frac{\sum_{c=1}^{N-1} u_{c j} \left(1 - p_{c j}^{\alpha-1} \right)}{\alpha - 1}$
Species equivalents ^c	$E = 1 / \sum_{i=1}^N \sum_{j=1}^N p_i p_j s_{ij}$	${}^{\alpha}E = \left(\sum_{j=1}^N p_j \left(\sum_{i=1}^N p_i s_{ij} \right)^{\alpha-1} \right)^{\frac{1}{(1-\alpha)}}$	${}^{\alpha}E^* = \left(\sum_{j=1}^N p_j \left(1 - \frac{\max_i(d_{ij})}{d_{\max}} + \sum_{c=1}^{N-1} \frac{u_{c j}}{d_{\max}} p_{c j}^{\alpha-1} \right) \right)^{\frac{1}{(1-\alpha)}}$



2.18

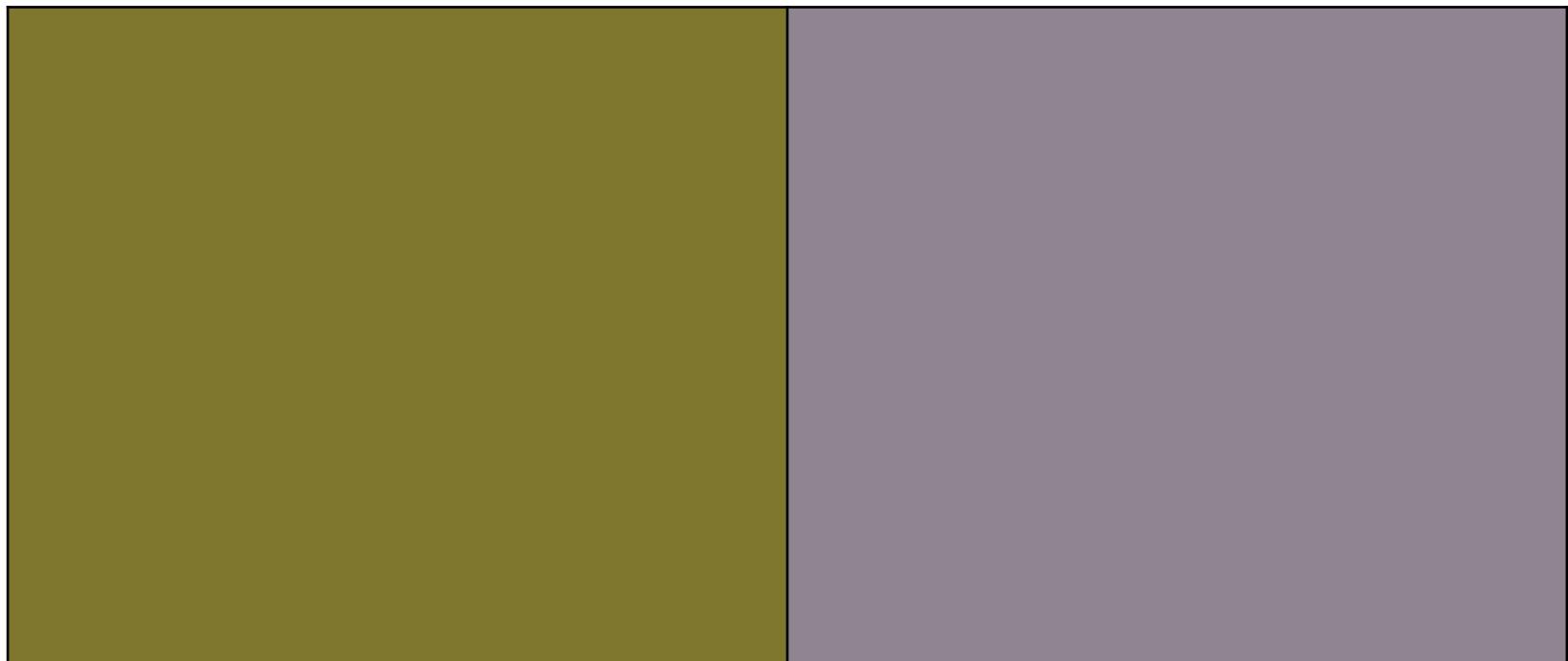


2.38

Range Size

Large

Small



Range Size

Large

Small

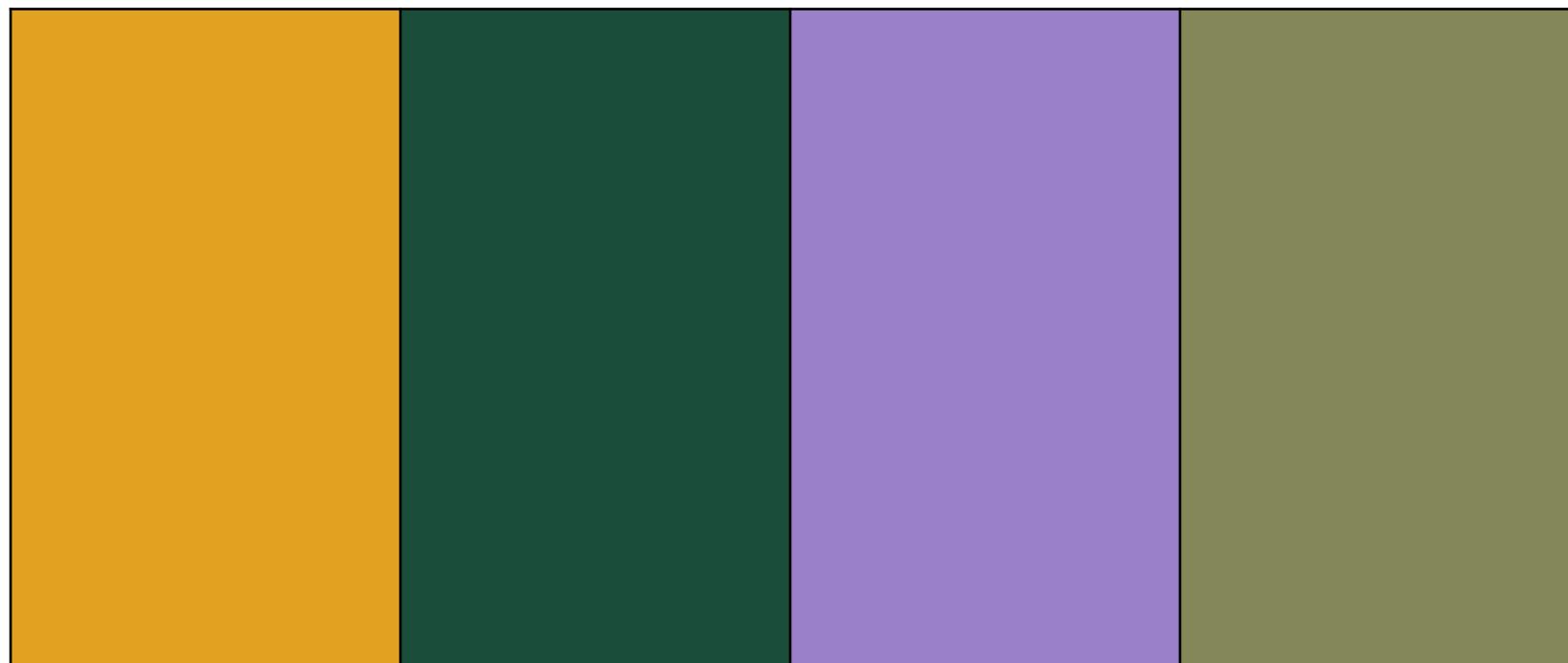
**Functional
Distinctiveness**

Distinct

Indistinct

Distinct

Indistinct



Range Size

Large

Small

**Functional
Distinctiveness**

Distinct

Indistinct

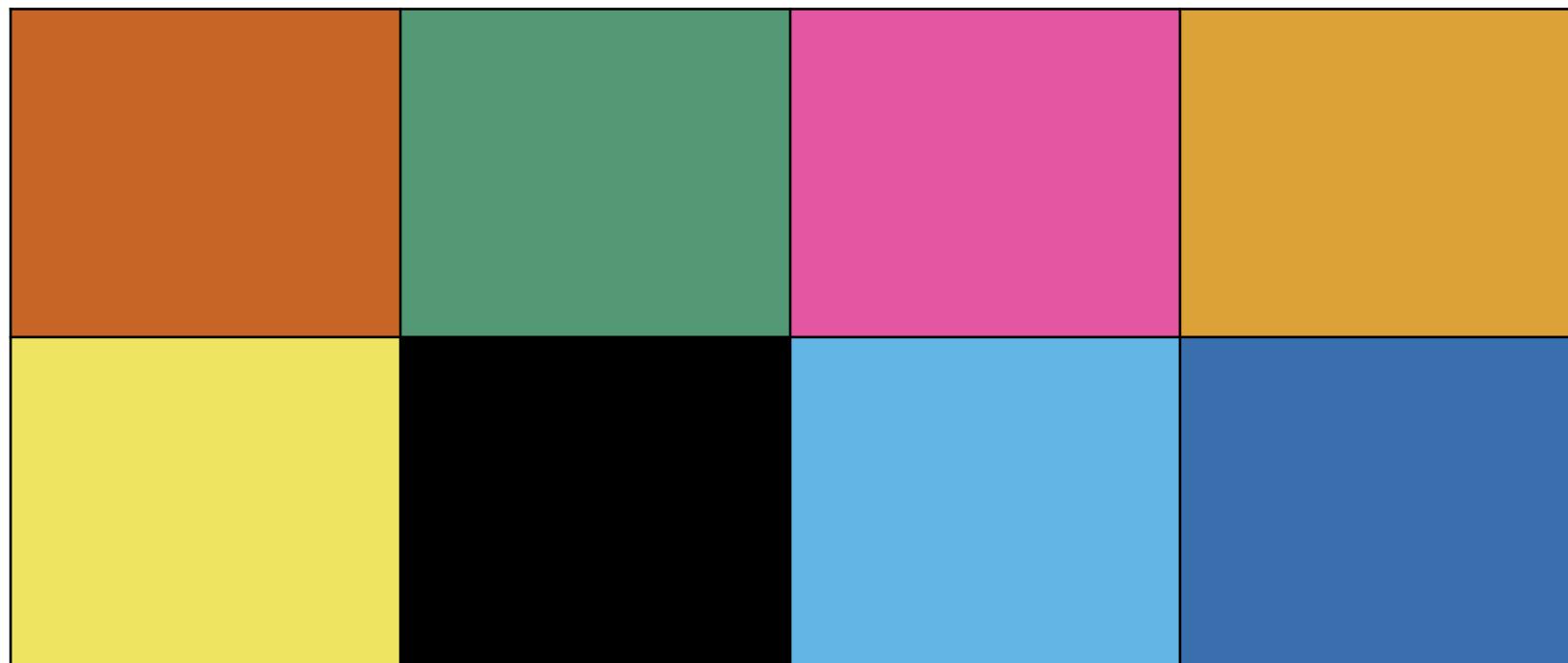
Distinct

Indistinct

**Evolutionary
Distinctiveness**

Distinct

Indistinct



Range Size

Large

Small

**Functional
Distinctiveness**

Distinct

Indistinct

Distinct

Indistinct

**Evolutionary
Distinctiveness**

Distinct

Indicator

Adaptable
Survivor

Classically Rare

Relict

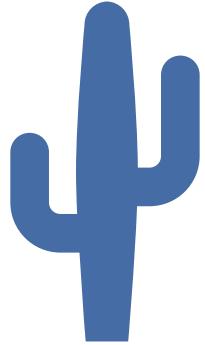
Indistinct

High Invasive
Potential

Common

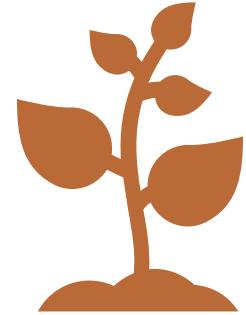
Endemic

Environmentally
Rare



Environmentally Rare

- High stress tolerance
- General mutualisms
- Recent speciation



Indicator

- Indicative of ecosystem health
- Disruptive outside of native range

What is the best definition of discrete rarity?

The word 'Abundance' is the central, largest word in the cloud. Surrounding it are various other terms related to rarity definitions, such as 'Quartiles', 'K Means Clustering', 'Mean Euclidean Distance', 'Minimum Euclidean Distance', 'Area of Occupancy', 'Trait Distinctiveness', 'Evolutionary Distinctiveness', 'Taxonomic Frequency', 'Trait Uniqueness', 'Range Size', 'Mean Pair-Wise Distance', 'Extent of Occurrence', and 'Percentiles'. These terms are arranged in a roughly circular pattern around the central 'Abundance' word.

Quartiles
K Means Clustering
Mean Euclidean Distance
Minimum Euclidean Distance
Area of Occupancy

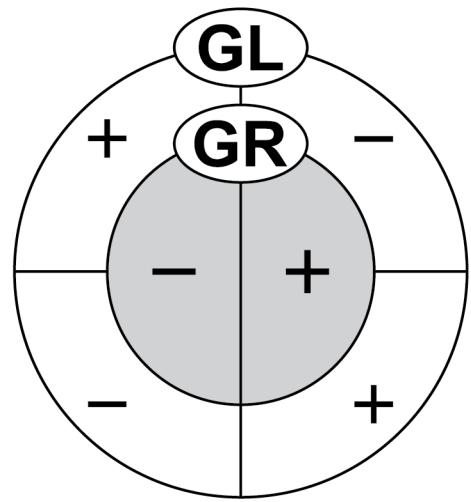
Abundance

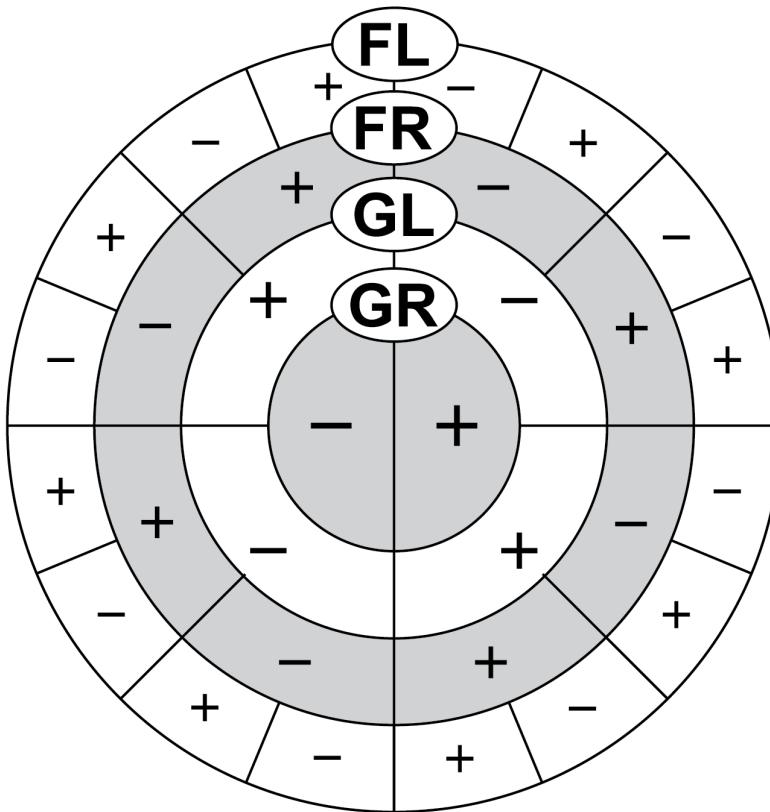
Trait Distinctiveness
Evolutionary Distinctiveness
Taxonomic Frequency

Trait Uniqueness

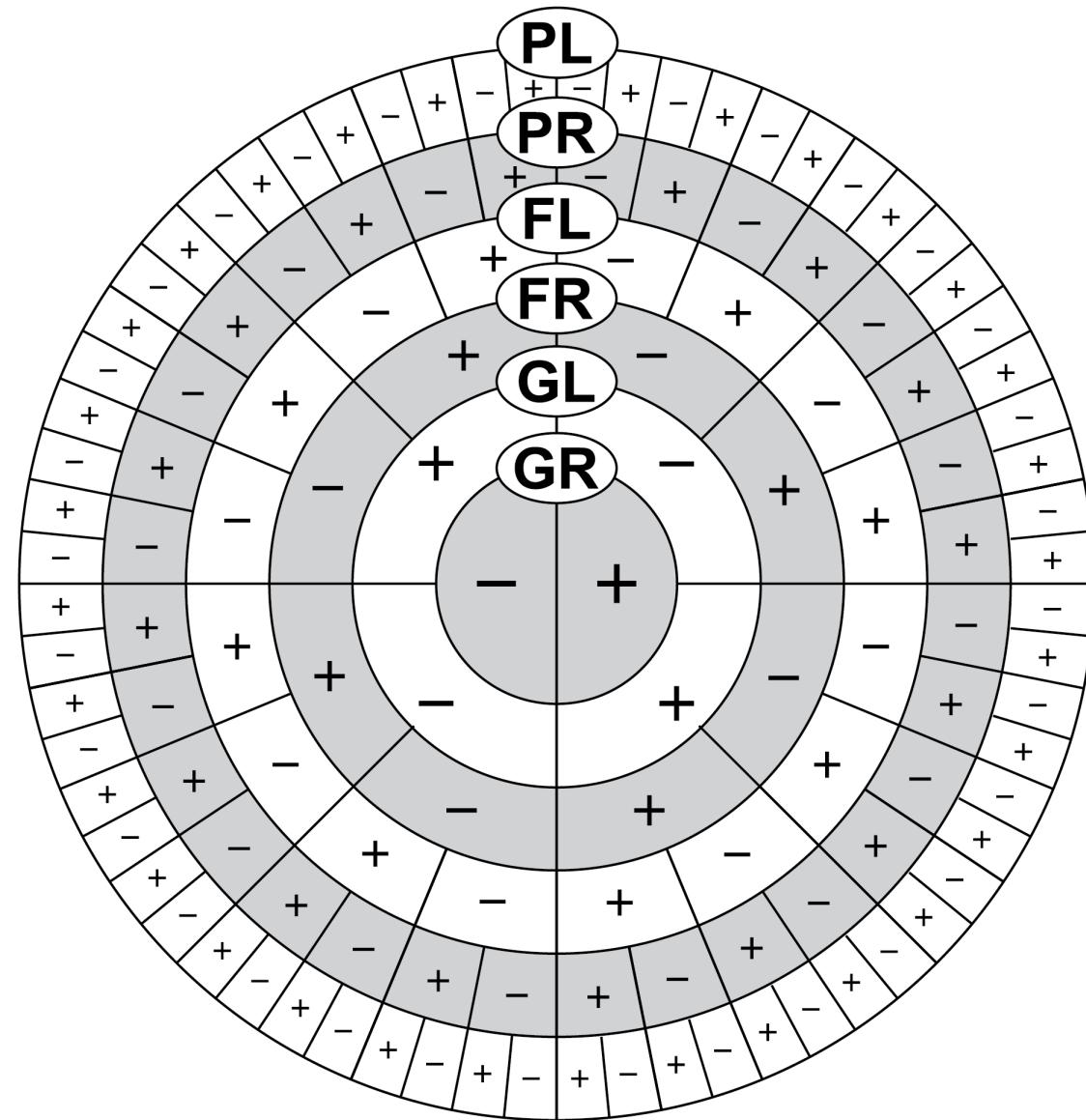
Range Size

Mean Pair-Wise Distance
Extent of Occurrence
Percentiles

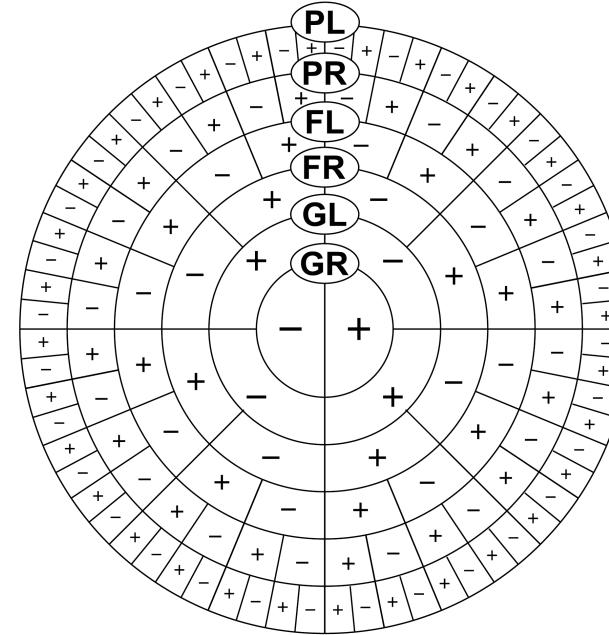




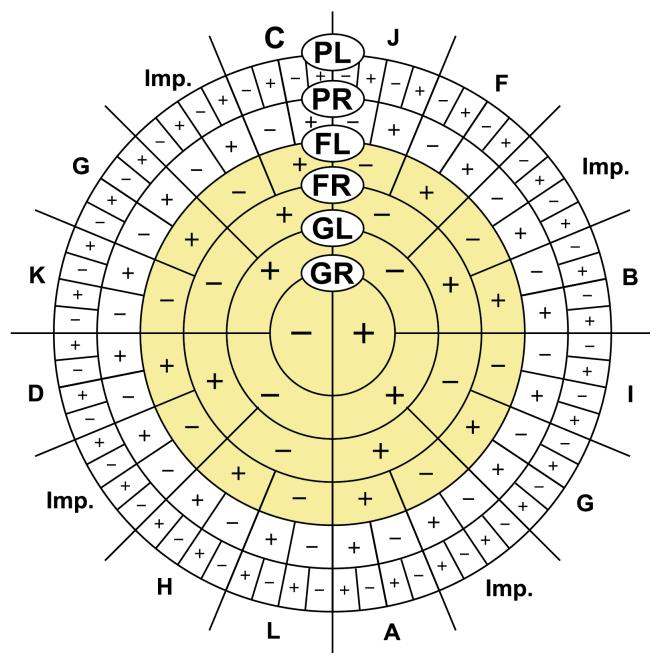
The Global Model of Discretized Rarity (GDR)



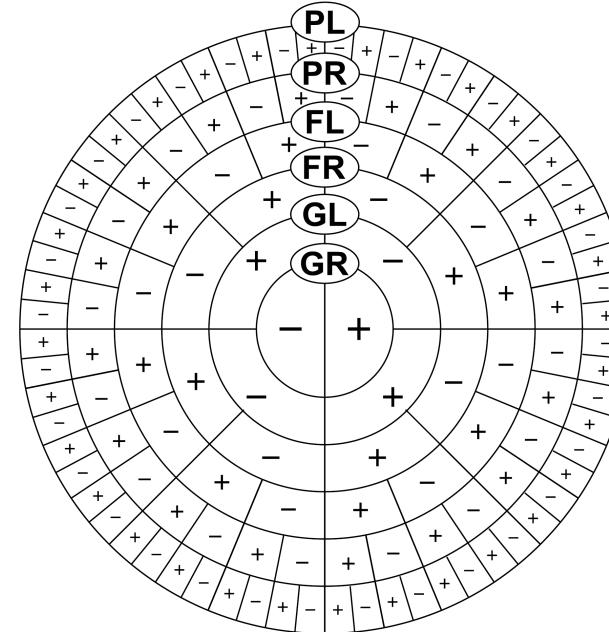
Global Model of Discretized Rarity



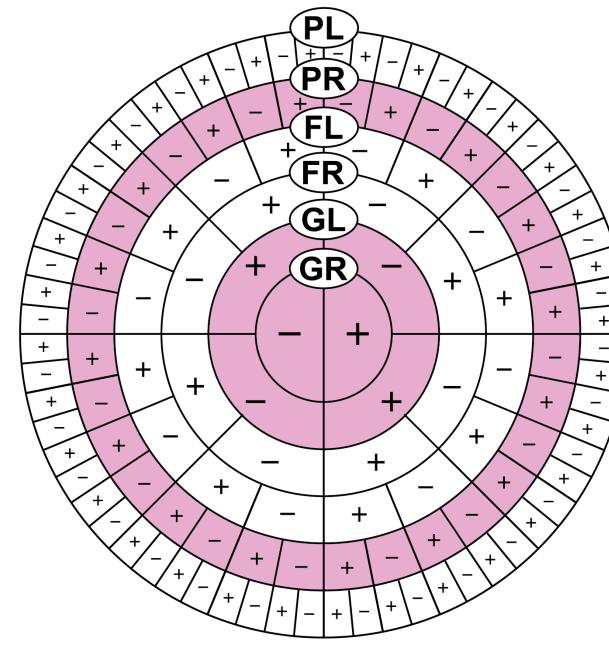
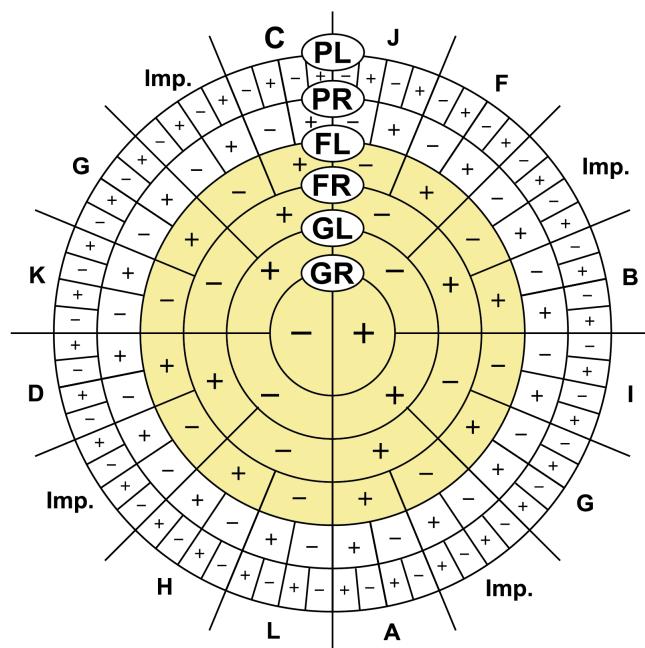
Functional Rarity Framework



Global Model of Discretized Rarity

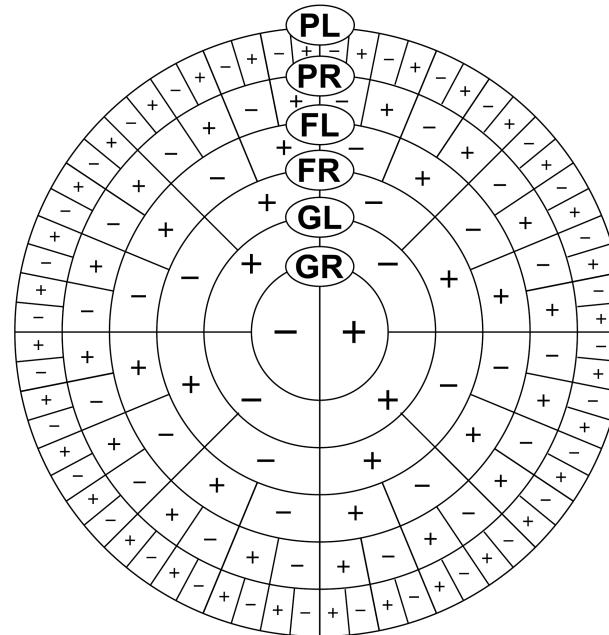


Functional Rarity Framework

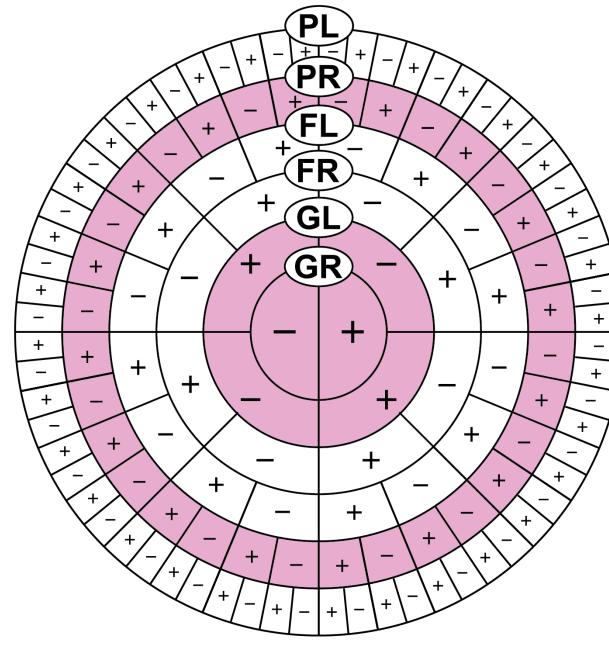
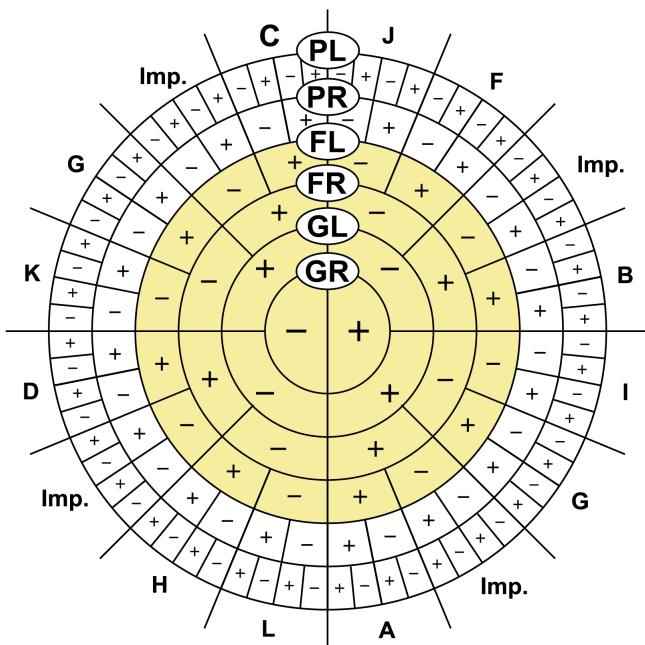


EDGE Protocol

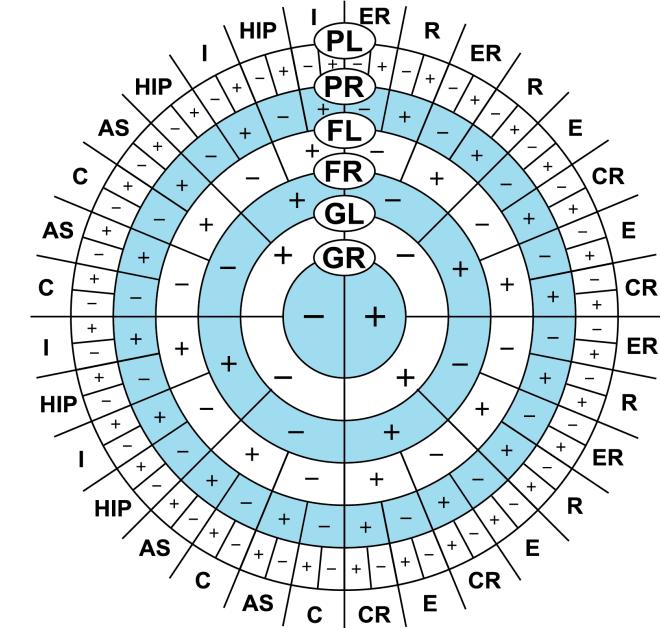
Global Model of Discretized Rarity



Functional Rarity Framework



Eco-Evolutionary Rarity



EDGE Protocol

Exploring the restrictions of GDR

1

Can we use rarity as a
tool to understand
ecological processes?

How?

2

Are rarity types robust
and ecologically
relevant?

Guide the selection and application of GDR restrictions



Data availability



Research/Management
Objectives



Methods & ability to explain
biological processes



Building a working global model



Building a working global model

Regional

GR = Extent of occurrence or taxonomic frequency

FR = Minimum or mean Euclidean distance

PR = Evolutionary Distinctiveness



Local

GL = Abundance

FL = Mean abundance weighted distance

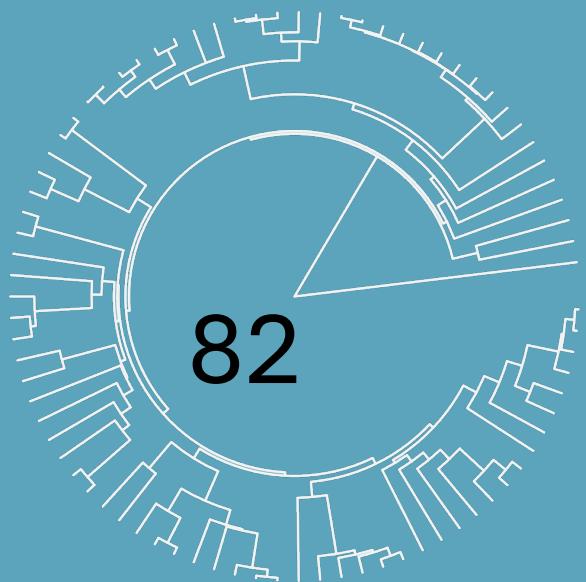
PL = Phylogenetic abundance weighted or unweighted mean pair-wise distance

British Plant Species

- Rabinowitz rarity was classified for 177 British flora in 1986.

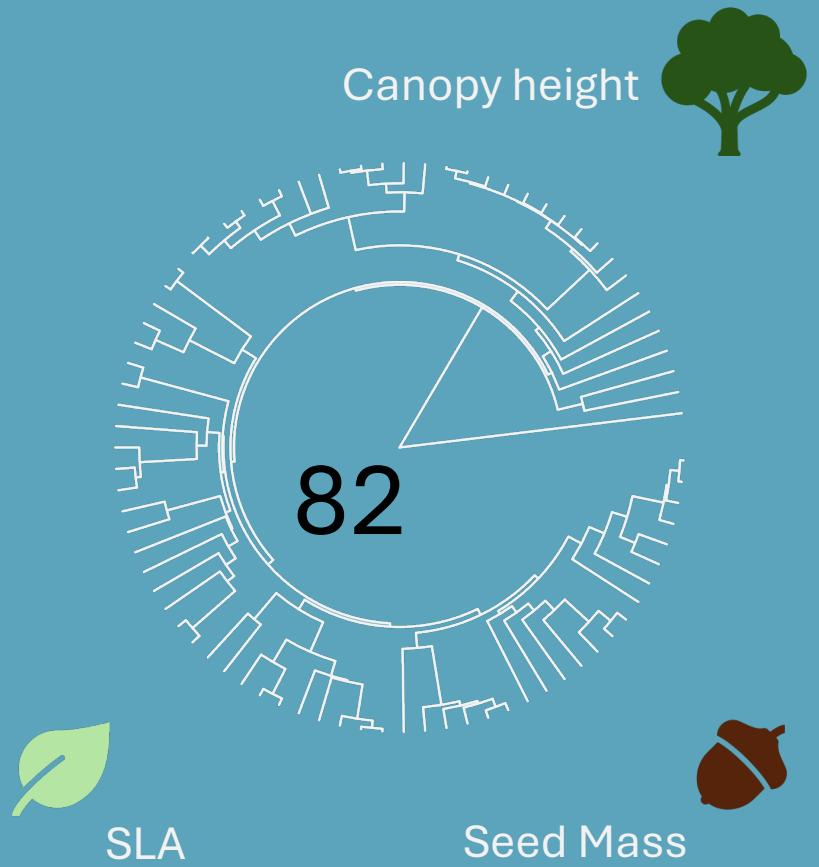
British Plant Species

- Rabinowitz rarity was classified for 177 British flora in 1986.



British Plant Species

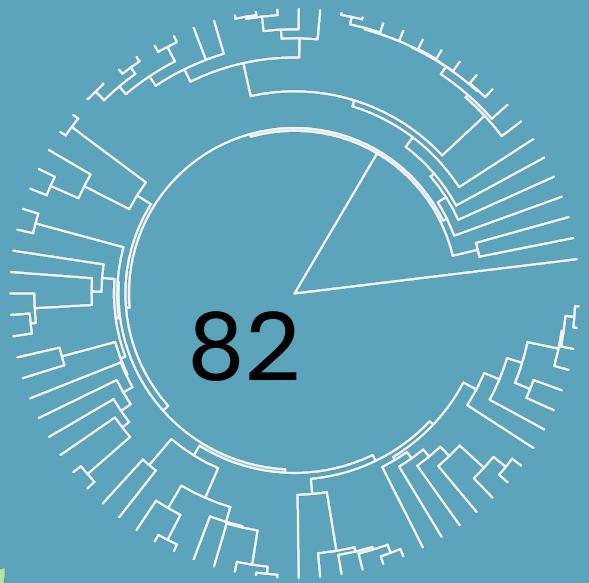
- Rabinowitz rarity was classified for 177 British flora in 1986.



British Plant Species

- Rabinowitz rarity was classified for 177 British flora in 1986.

Canopy height

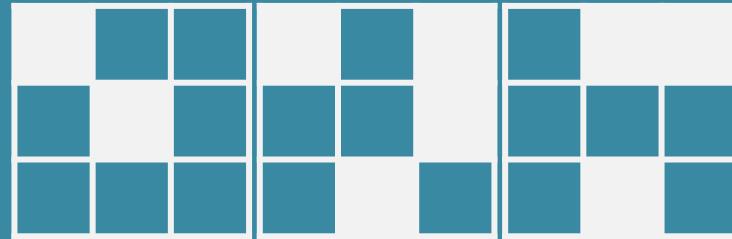


SLA



Seed Mass

Plant Atlas 2020



1970-1986

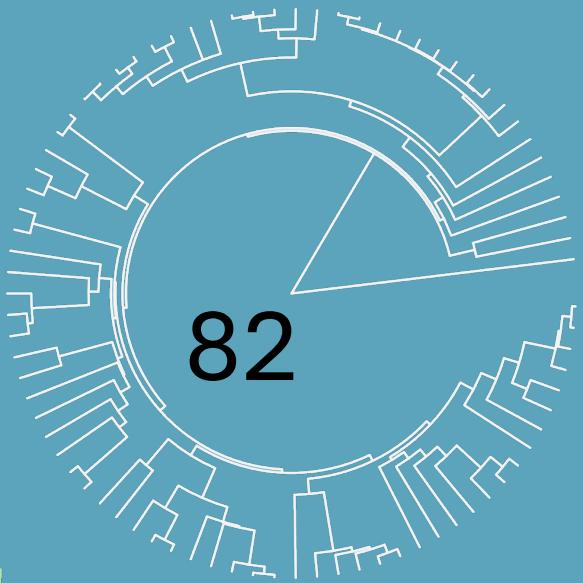
1987-1999

2000-2019

British Plant Species

- Rabinowitz rarity was classified for 177 British flora in 1986.

Canopy height

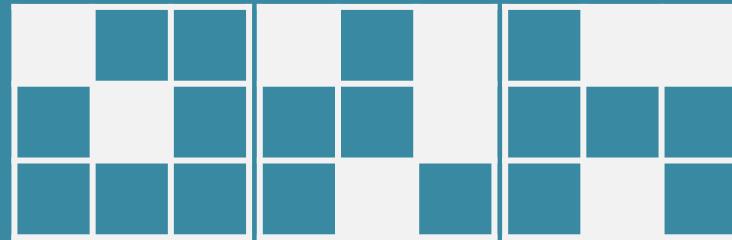


SLA



Seed Mass

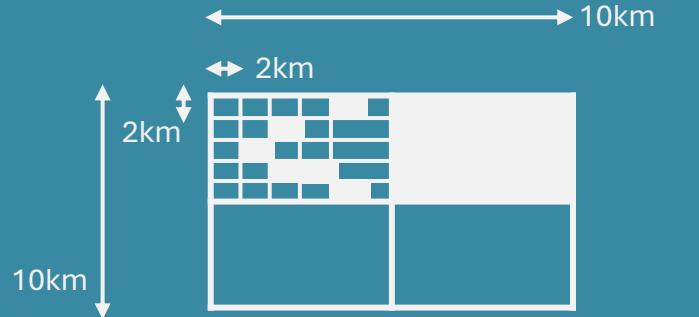
Plant Atlas 2020



1970-1986

1987-1999

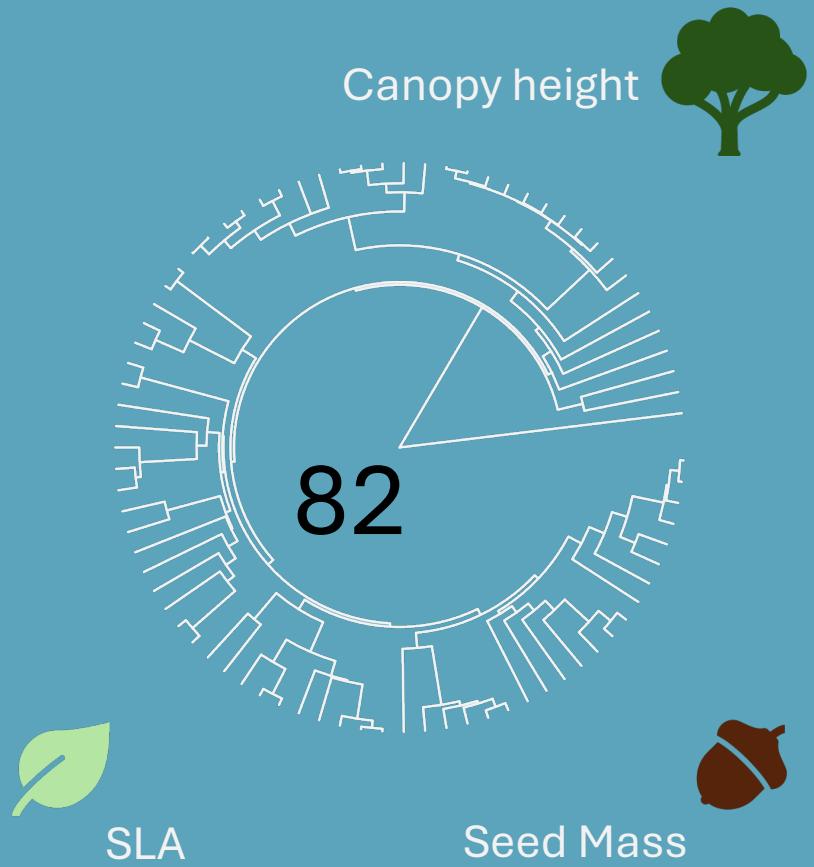
2000-2019



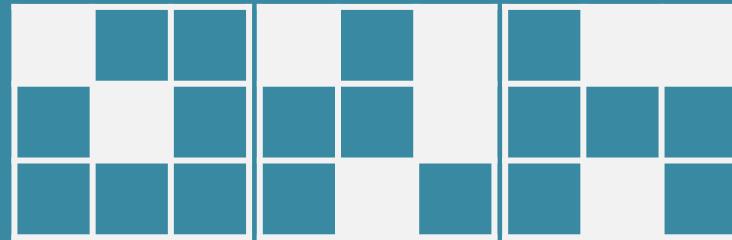
1970-1986

British Plant Species

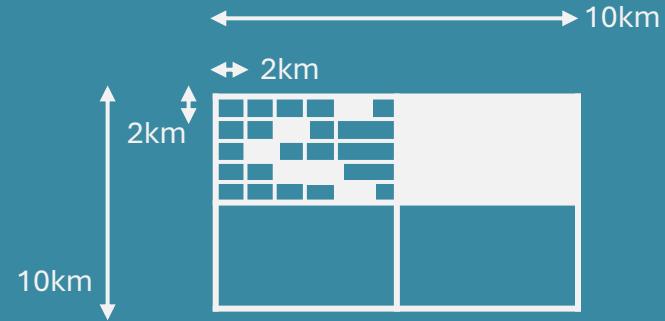
- Rabinowitz rarity was classified for 177 British flora in 1986.



Plant Atlas 2020



1970-1986 1987-1999 2000-2019

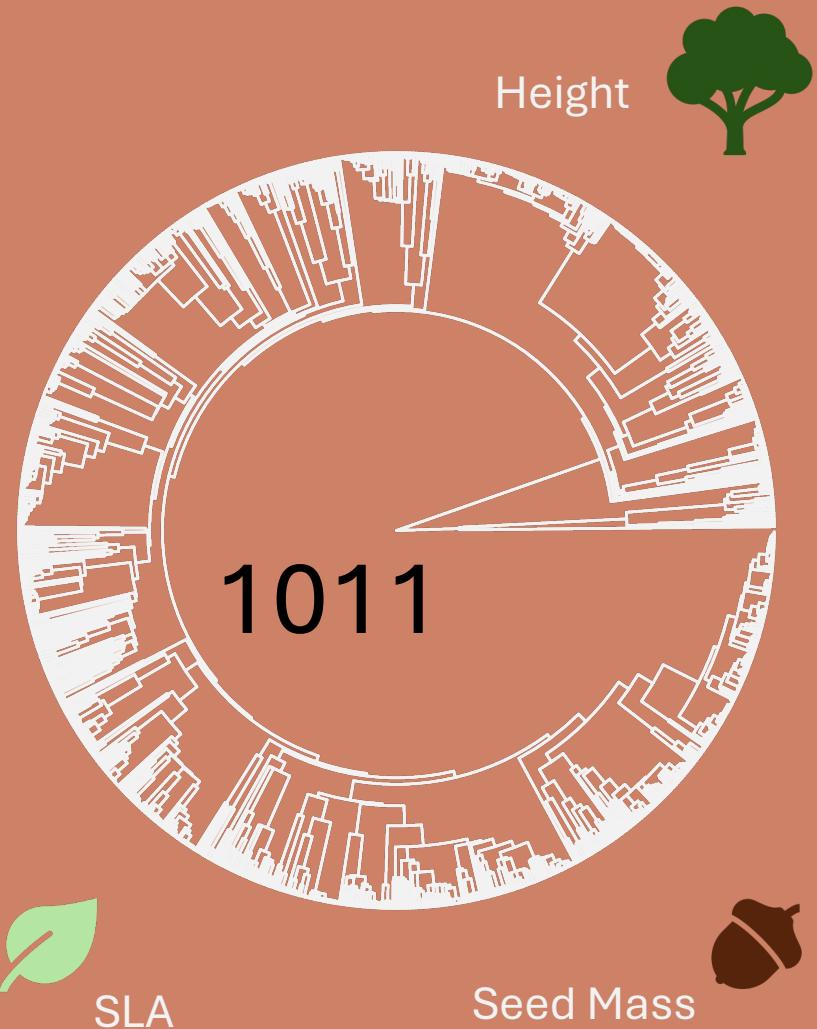


1970-1986

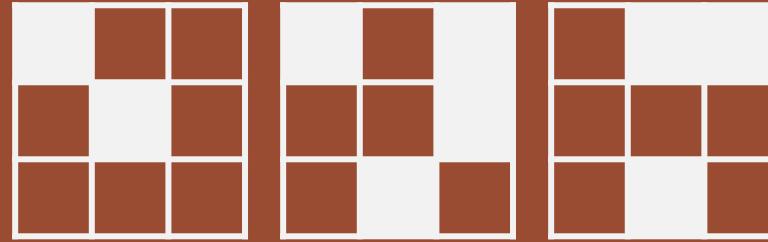


Flowering Window

British Plant Species

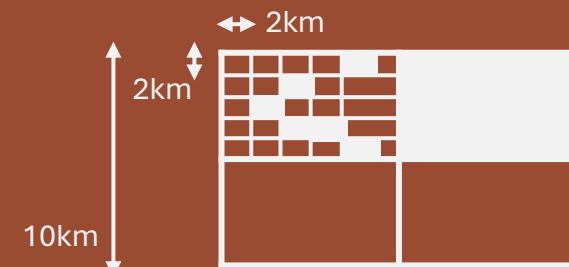


Plant Atlas 2020



1970-1986 1987-1999 2000-2019

10km 2km 10km

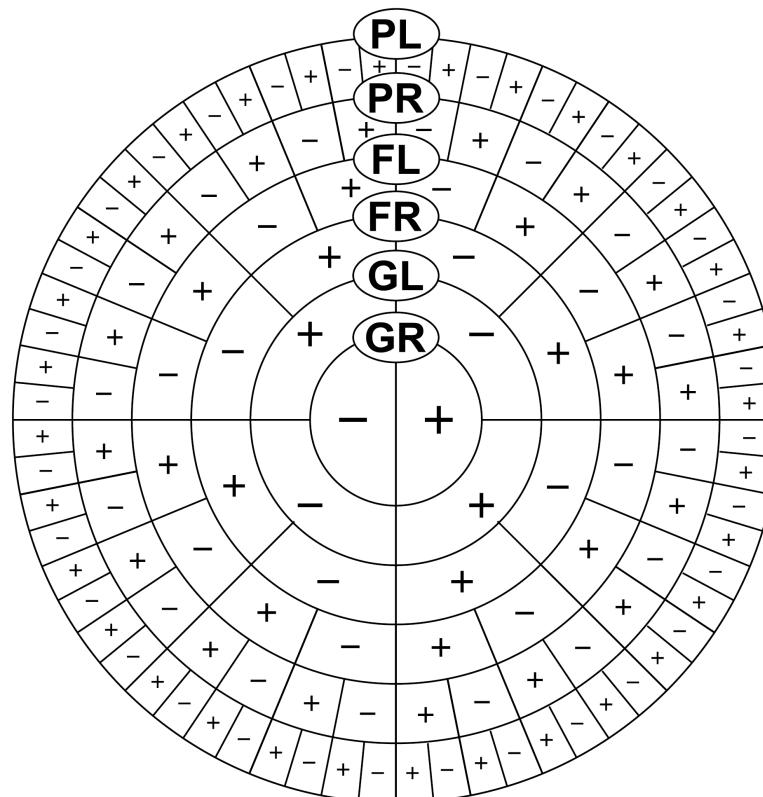


Flowering Window

82 Species

1011 Species

4 Thresholds X 2 GR X 2 FR X 2 PL = 32 Methods
63 Restrictions



82 Species

1011 Species

4 Thresholds X 2 GR X 2 FR X 2 PL = 32 Methods
63 Restrictions

For each method:

Distribution Change ~ Restriction

82 Species

1011 Species

4 Thresholds X 2 GR X 2 FR X 2 PL = 32 Methods
63 Restrictions

For each method:

Distribution Change ~ Restriction

Flowering Phenology ~ Restriction

82 Species

1011 Species

4 Thresholds X 2 GR X 2 FR X 2 PL = 32 Methods
63 Restrictions

For each method:

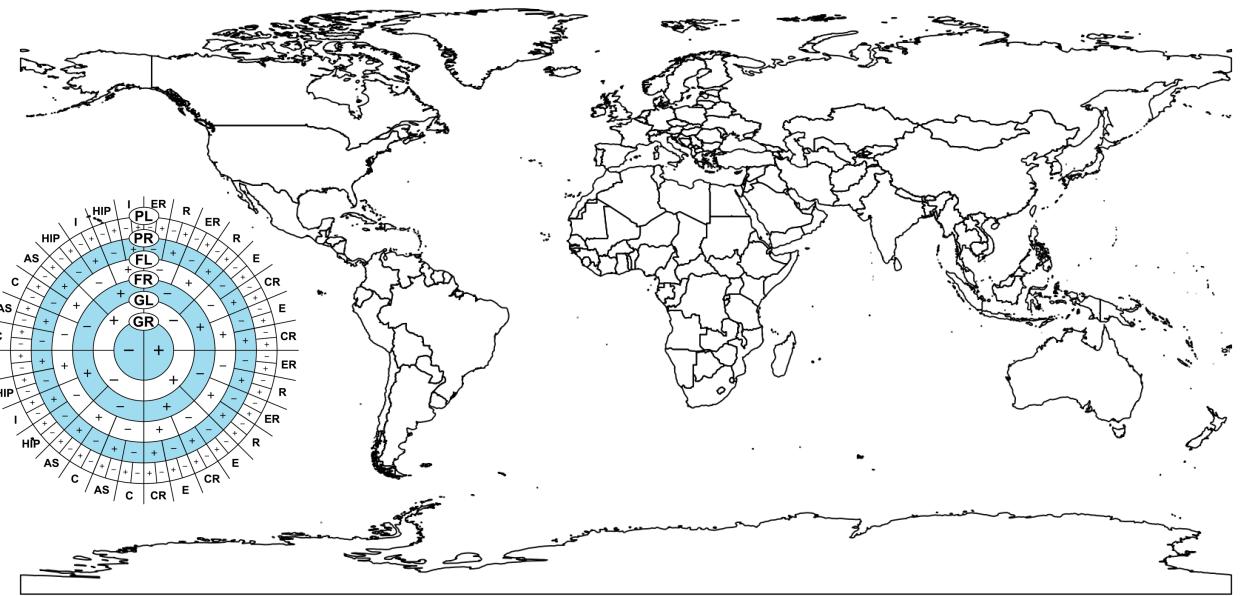
Distribution Change ~ Restriction

Flowering Phenology ~ Restriction

Best Method per Restriction

&

Best Restriction per Ecological Process

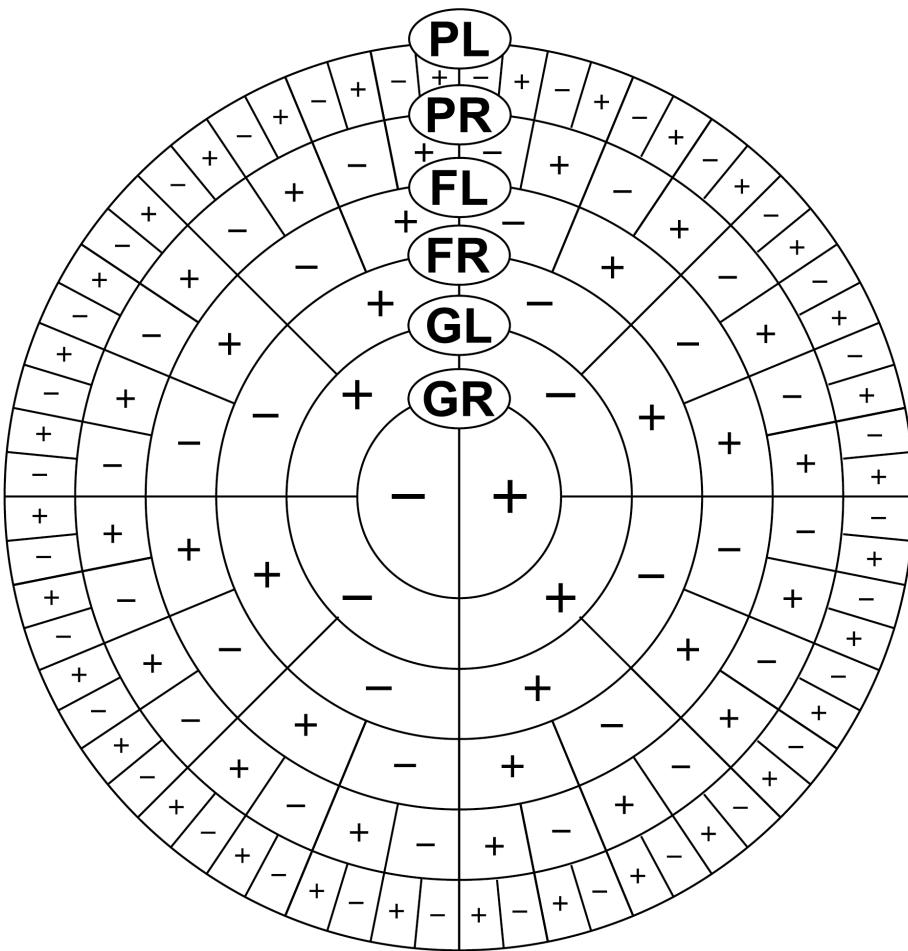


GRIIS
GLOBAL REGISTER OF
INTRODUCED AND INVASIVE SPECIES

Eco-Evolutionary Rarity of 5,611 Global Angiosperms

The Global Model of Discretized Rarity performed well across sample sizes...

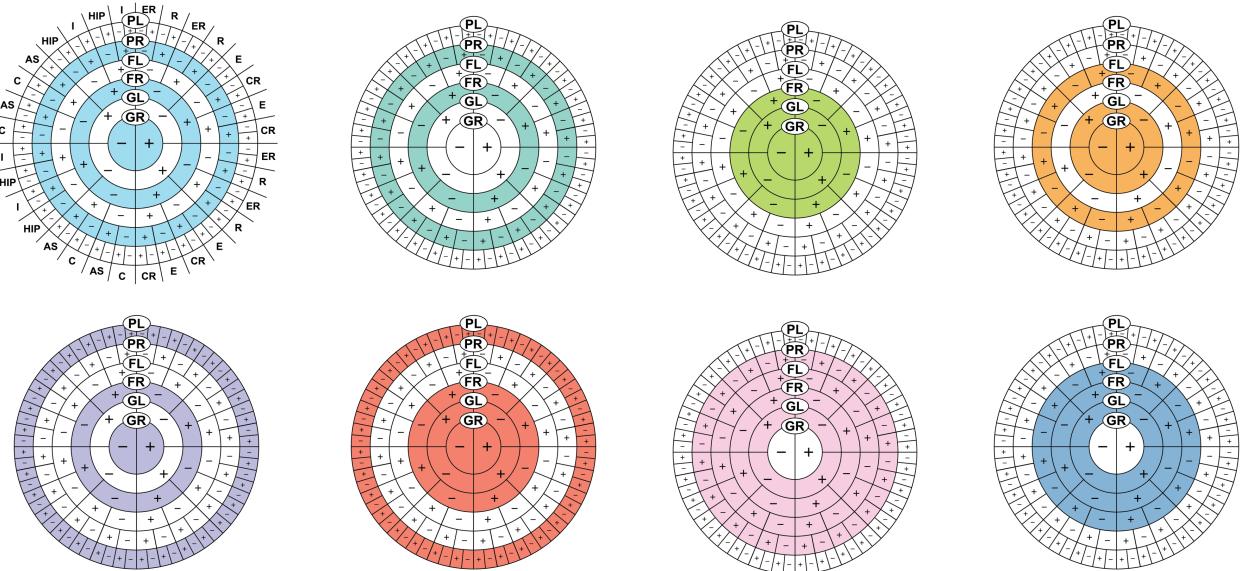
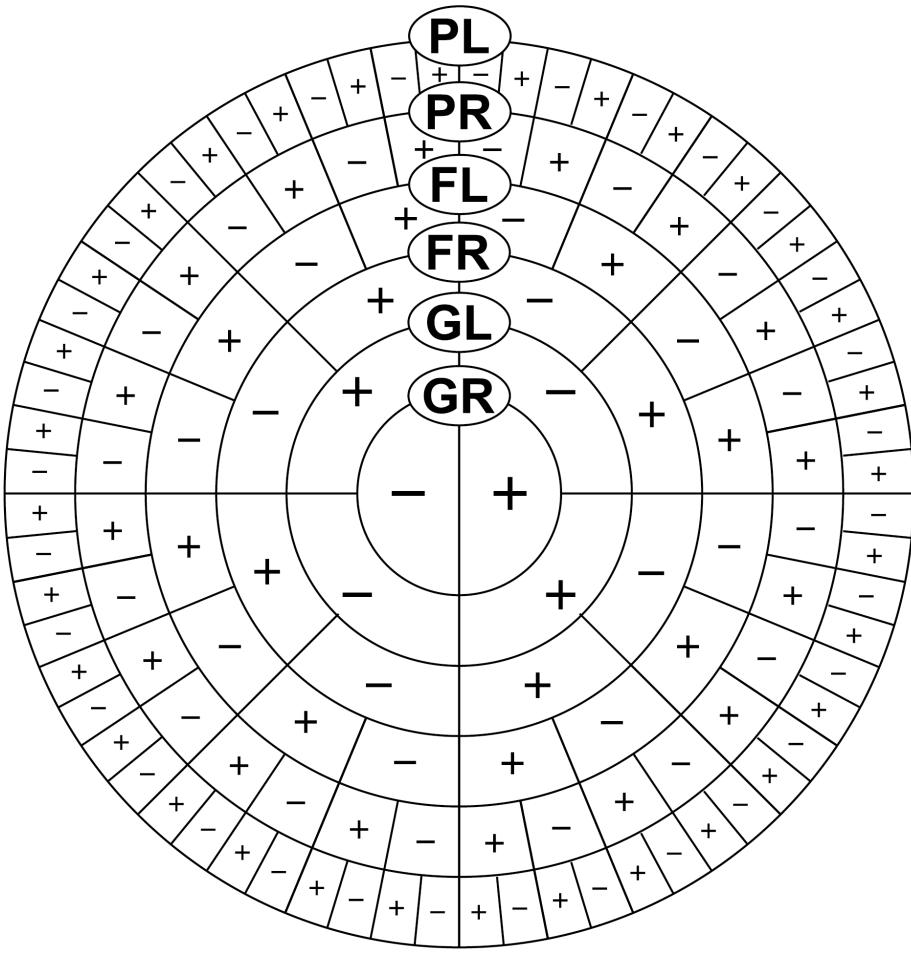
- 12.5% - 39% of variation in distribution change
- 13% - 49.1% of variation in flowering phenology



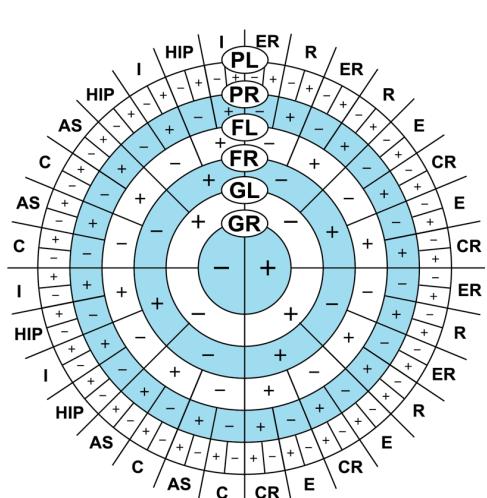
The Global Model of Discretized Rarity performed well across sample sizes...

- 12.5% - 39% of variation in distribution change
- 13% - 49.1% of variation in flowering phenology

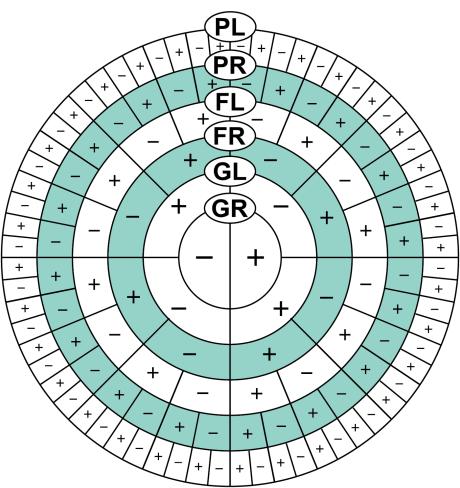
...but many restrictions outperformed the GDR



Small Sample Size



EER



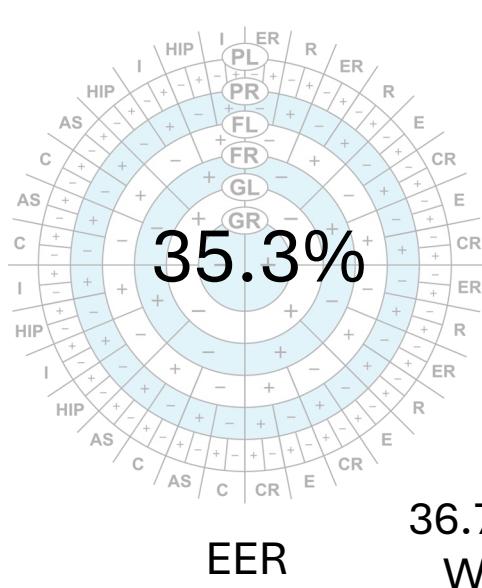
$F_R P_R$

Large Sample Size

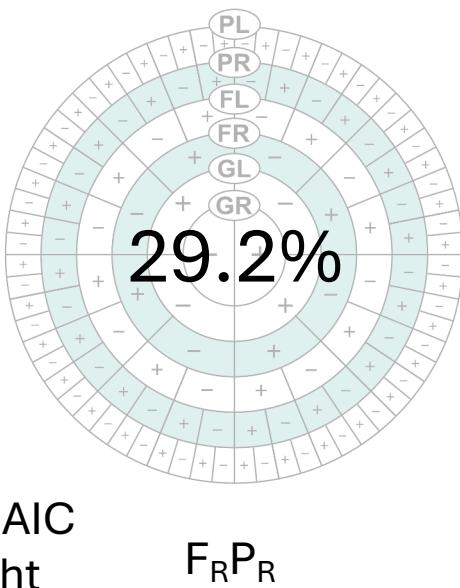
Distribution
Change

Flowering
Phenology

Small Sample Size



36.7% AIC
Weight

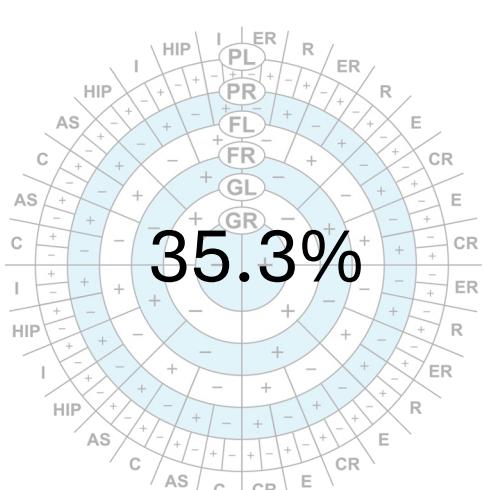


Large Sample Size

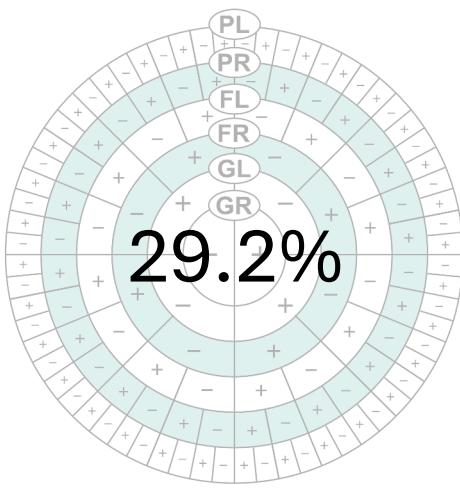
Distribution
Change

Flowering
Phenology

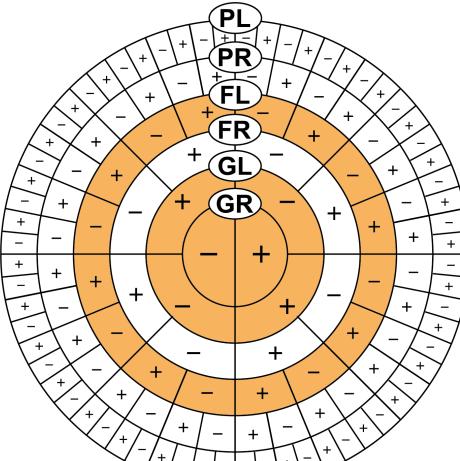
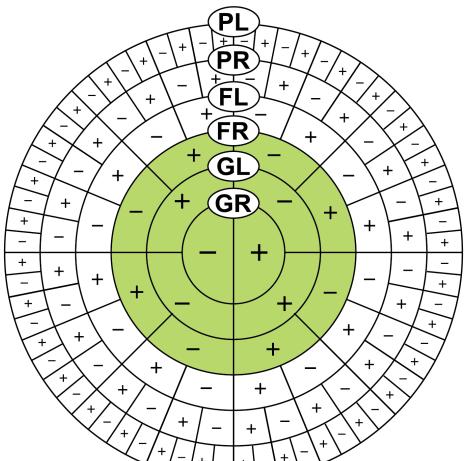
Small Sample Size



EER
36.7% AIC
Weight



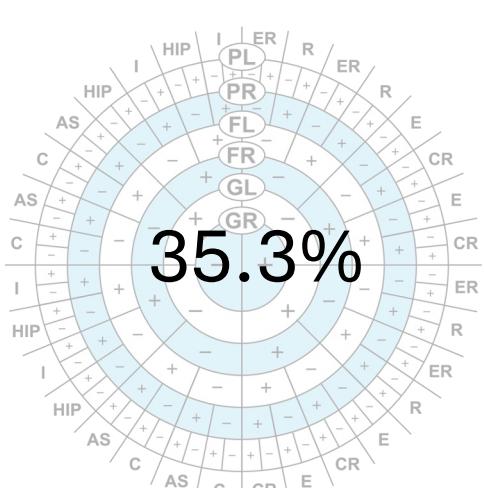
Distribution
Change



Flowering
Phenology

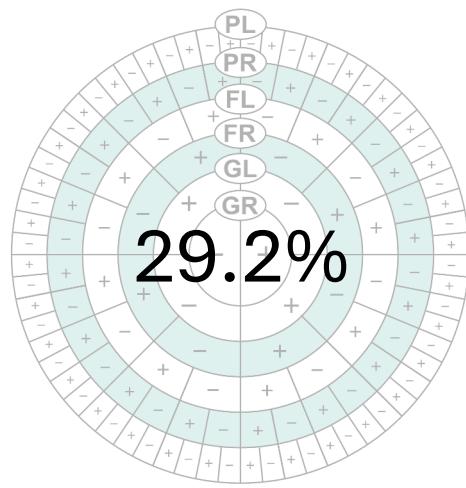
Large Sample Size

Small Sample Size



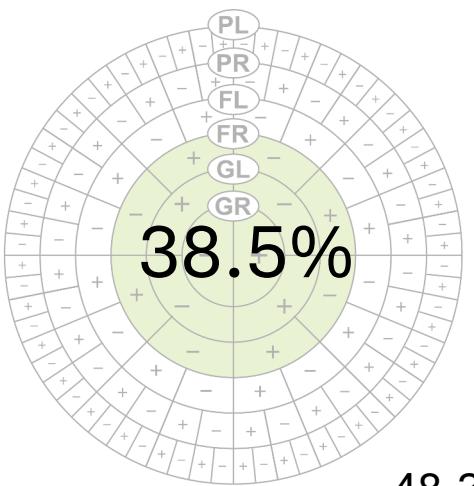
EER

36.7% AIC
Weight



29.2%

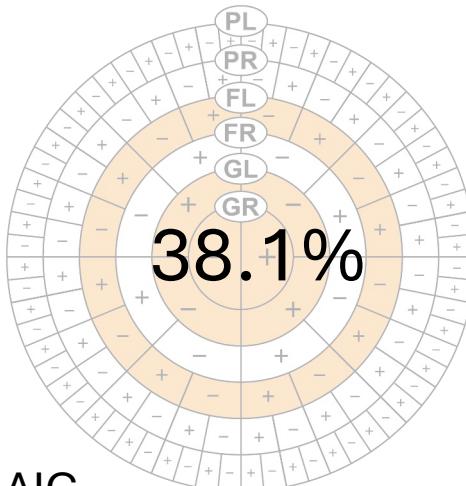
$F_R P_R$



38.5%

$G_{RL} F_R$

48.3% AIC
Weight



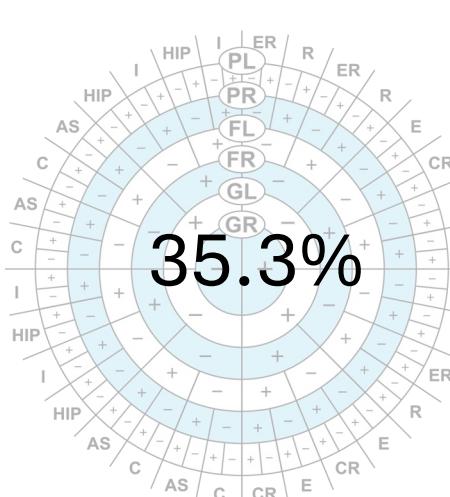
$G_{RL} F_L$

Large Sample Size

Distribution
Change

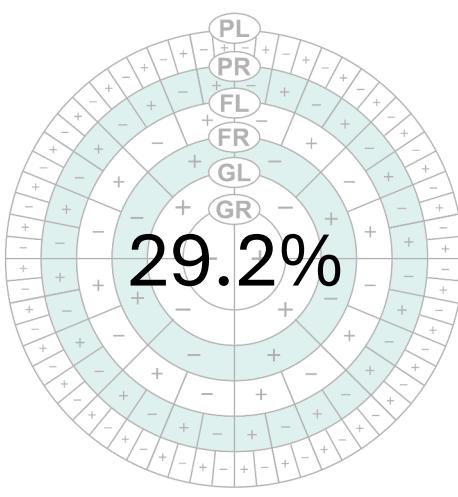
Flowering
Phenology

Small Sample Size



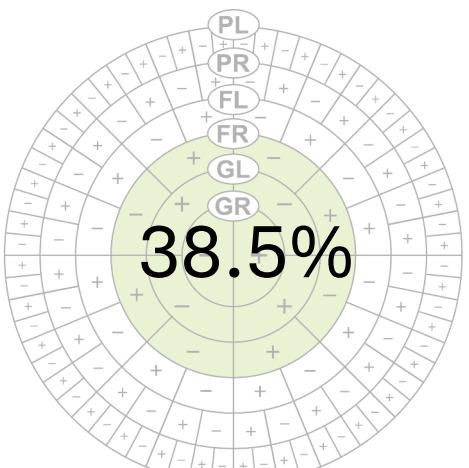
EER

36.7% AIC
Weight



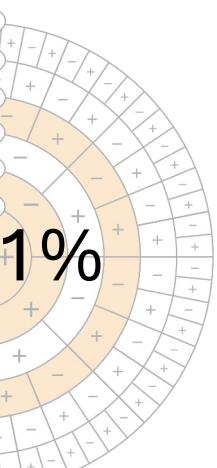
$F_R P_R$

Distribution
Change



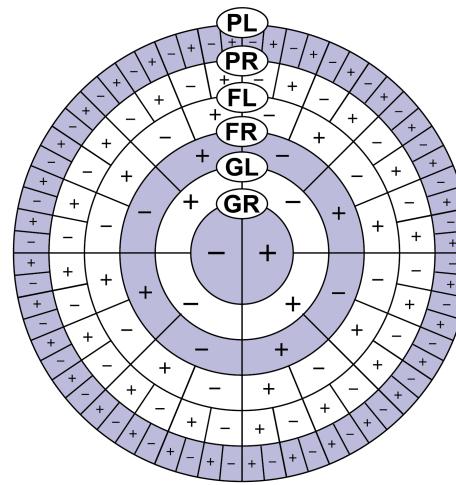
$G_{RL} F_R$

48.3% AIC
Weight

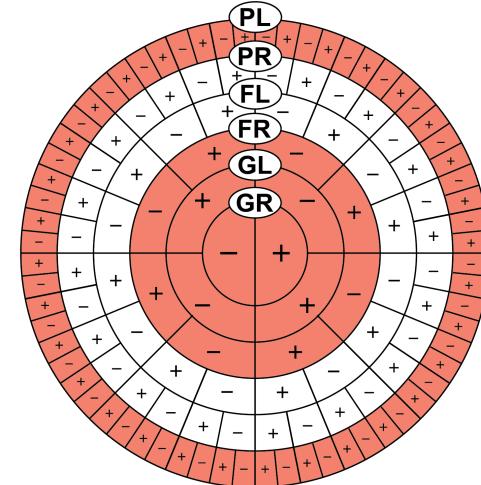


$G_{RL} F_L$

Large Sample Size



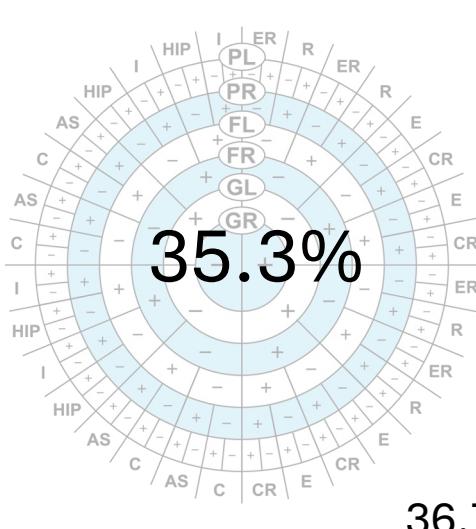
$G_R F_R P_L$



$G_{RL} F_R P_L$

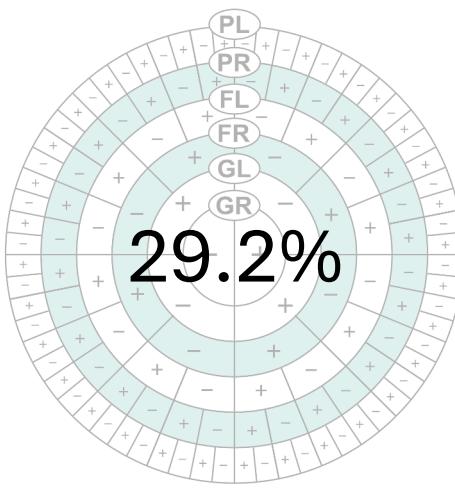
Flowering
Phenology

Small Sample Size



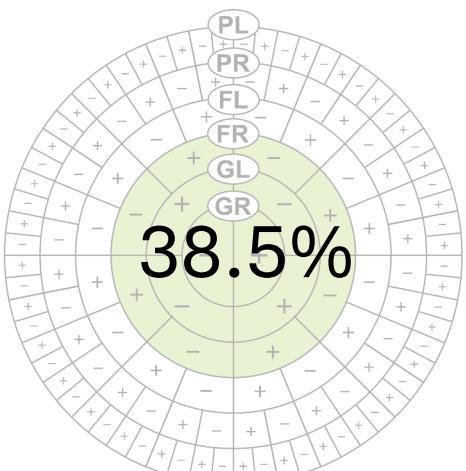
EER

36.7% AIC
Weight



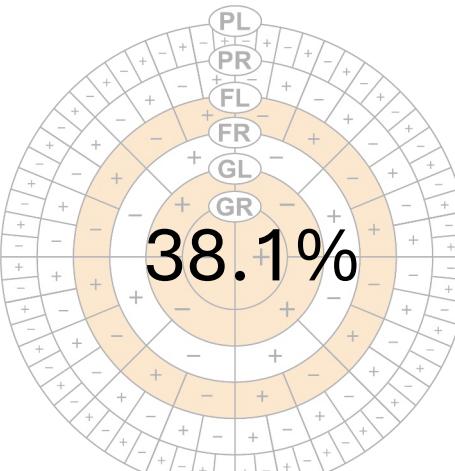
$F_R P_R$

Distribution
Change



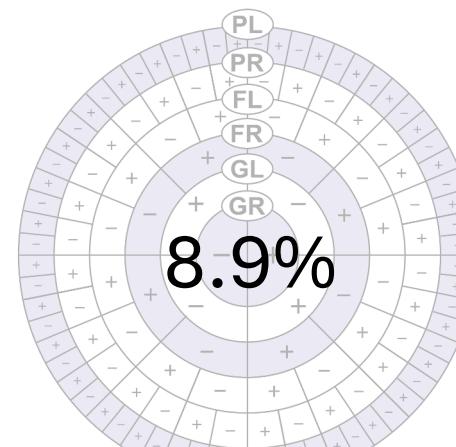
$G_{RL} F_R$

48.3% AIC
Weight



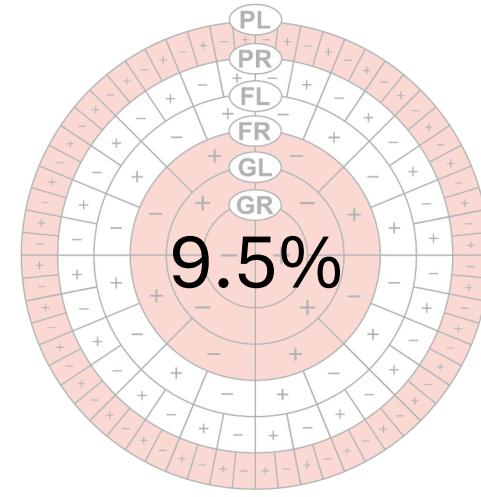
$G_{RL} F_L$

Large Sample Size



$G_R F_R P_L$

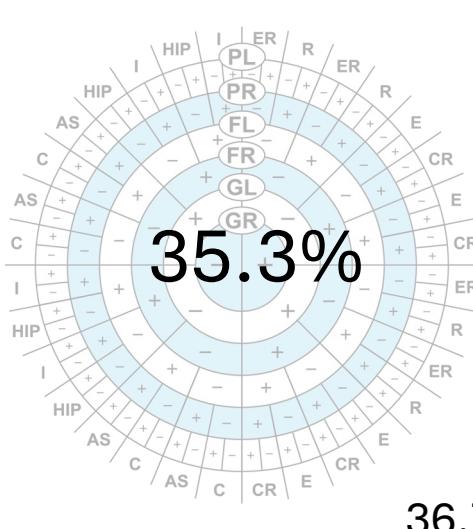
89.3% AIC
Weight



$G_{RL} F_R P_L$

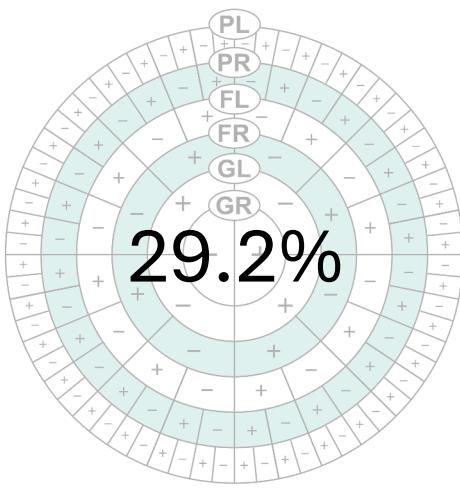
Flowering
Phenology

Small Sample Size



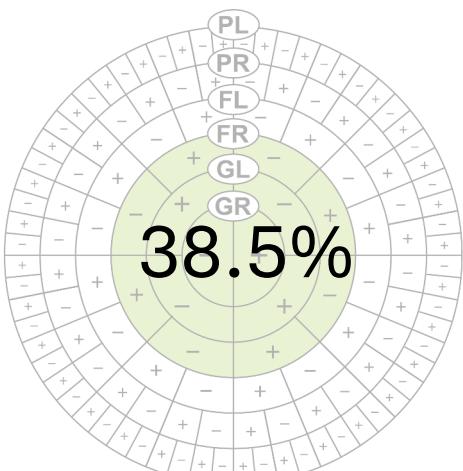
EER

36.7% AIC
Weight



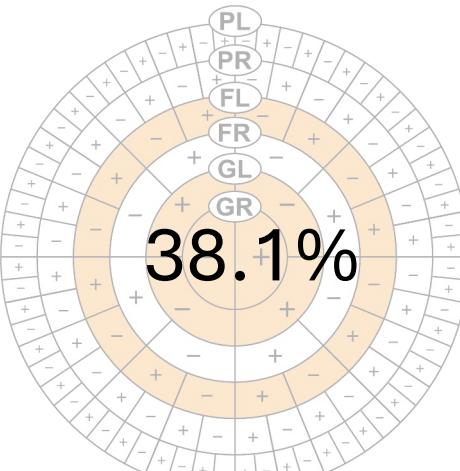
$F_R P_R$

Distribution
Change



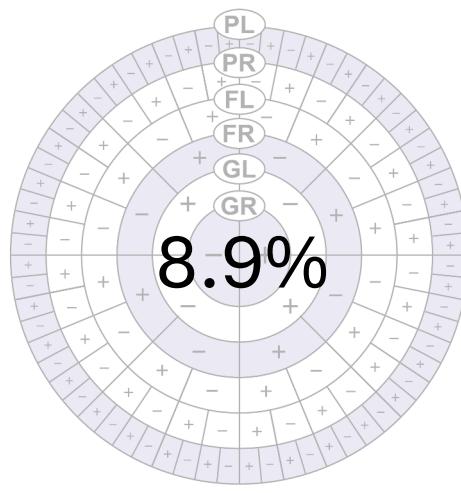
$G_{RL} F_R$

48.3% AIC
Weight



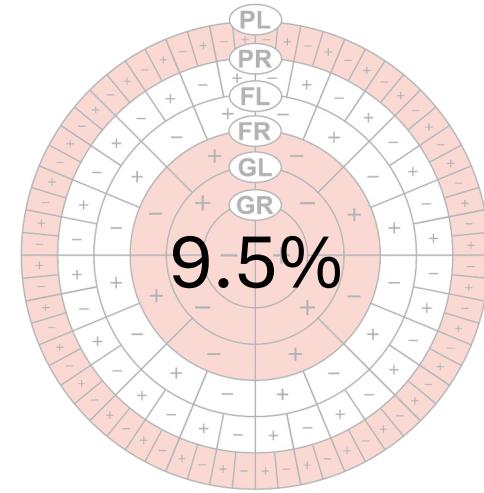
$G_{RL} F_L$

Large Sample Size



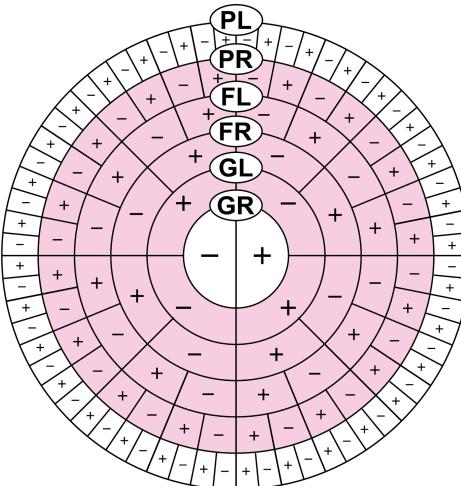
$G_R F_R P_L$

89.3% AIC
Weight

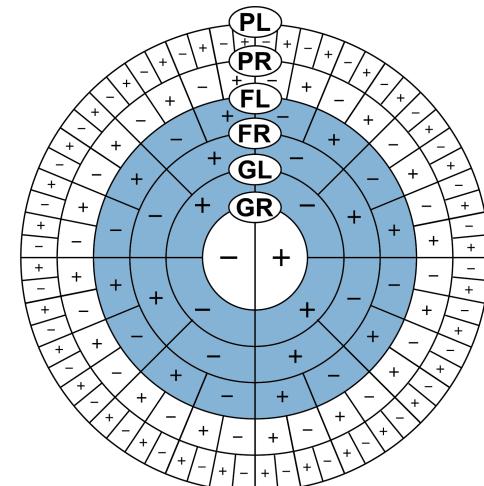


$G_{RL} F_R P_L$

Flowering
Phenology

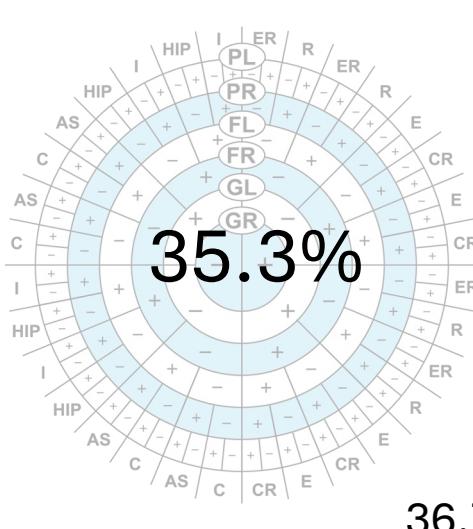


$G_L F_{RL} P_R$

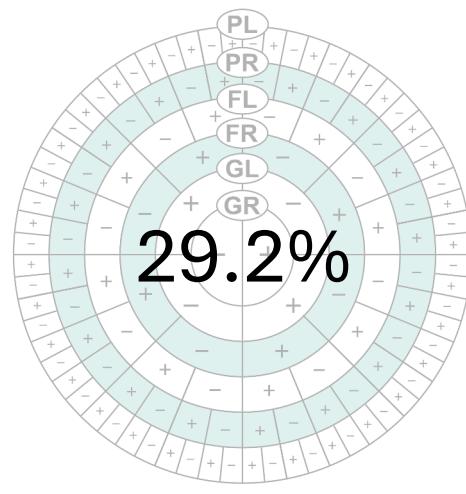


$G_L F_{RL}$

Small Sample Size

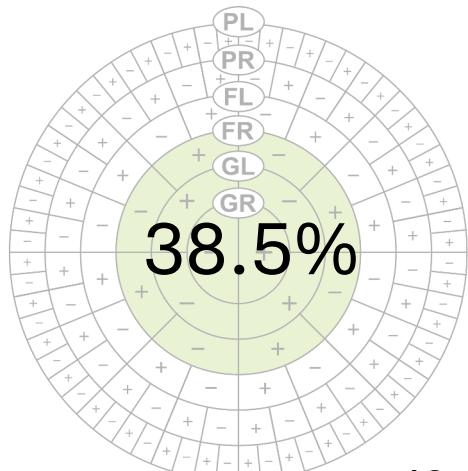


EER



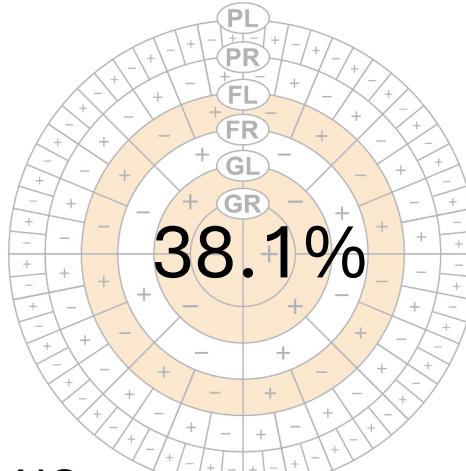
36.7% AIC
Weight

$F_R P_R$



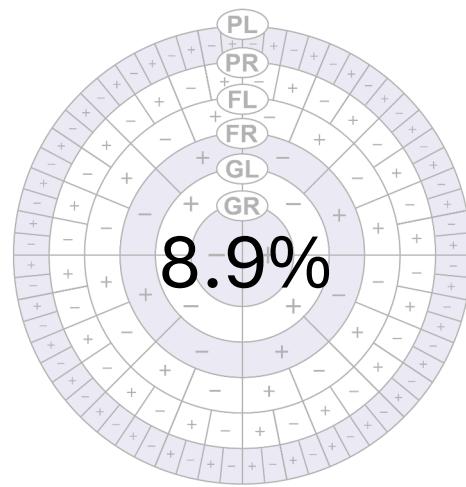
$G_{RL} F_R$

48.3% AIC
Weight



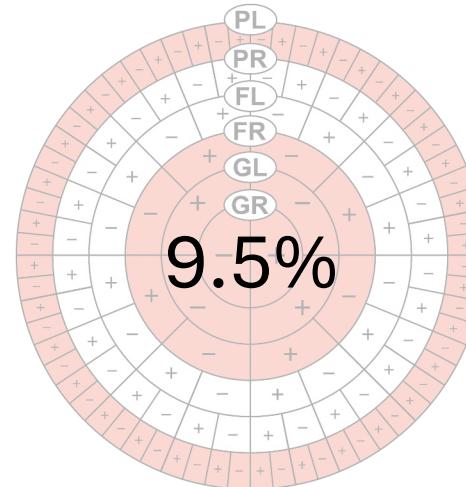
$G_{RL} F_L$

Large Sample Size



$G_R F_R P_L$

89.3% AIC
Weight



$G_{RL} F_R P_L$

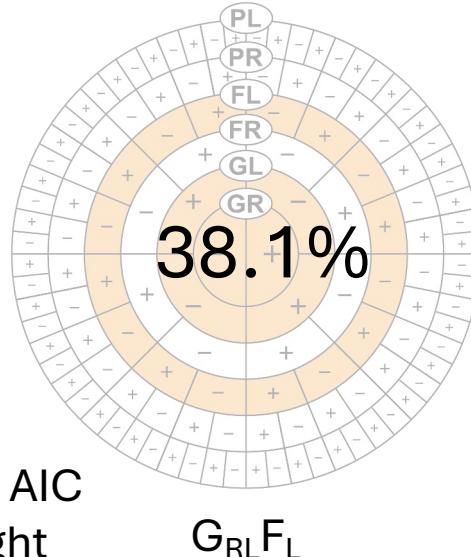
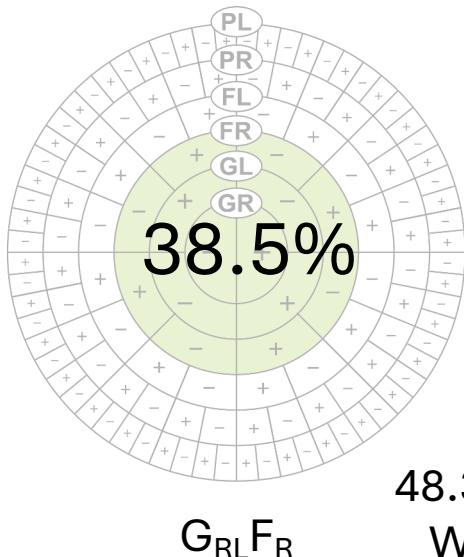
Distribution
Change

Flowering
Phenology

Small Sample Size

Dimension	AIC Importance
PR	96.7%
FR	91.6%
GR	73.1%
FL	23.8%
PL	15.7%
GL	15.6%

Distribution
Change



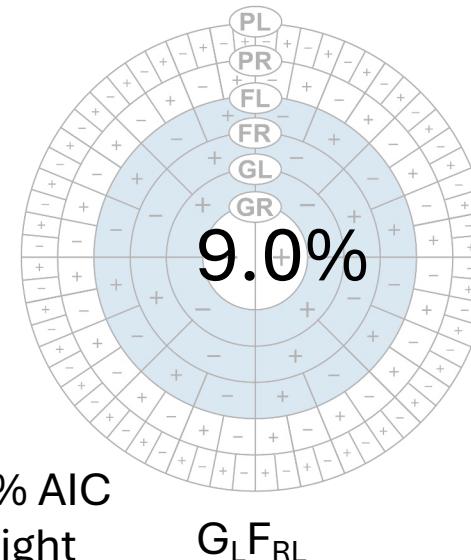
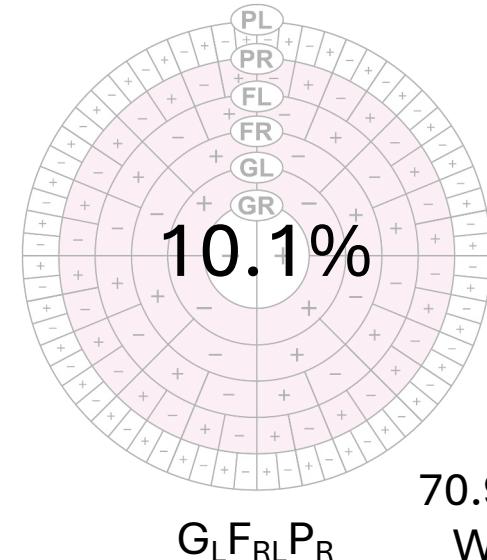
48.3% AIC
G_{RL}F_R Weight

G_{RL}F_L

Large Sample Size

Dimension	AIC Importance
GR	100%
FR	99.8%
PL	99.3%
GL	16.6%
PR	9.2%
FL	1.1%

Flowering
Phenology



70.9% AIC
G_LF_{RL}P_R Weight

G_LF_{RL}

Small Sample Size

Dimension	AIC Importance
PR	96.7%
FR	91.6%
GR	73.1%
FL	23.8%
PL	15.7%
GL	15.6%

Large Sample Size

Dimension	AIC Importance
GR	100%
FR	99.8%
PL	99.3%
GL	16.6%
PR	9.2%
FL	1.1%

Dimension	AIC Importance
GR	100%
GL	100%
FR	69.8%
FL	45.1%
PR	28.7%
PL	18.5%

Distribution Change

Flowering Phenology

Dimension	AIC Importance
GL	100%
FR	100%
FL	99.6%
PR	56.3%
PL	23.9%
GR	8.0%

Small Sample Size

Dimension	AIC Importance
PR	96.7%
FR	91.6%
GR	73.1%
FL	23.8%
PL	15.7%
GL	15.6%

Large Sample Size

Dimension	AIC Importance
GR	100%
FR	99.8%
PL	99.3%
GL	16.6%
PR	9.2%
FL	1.1%

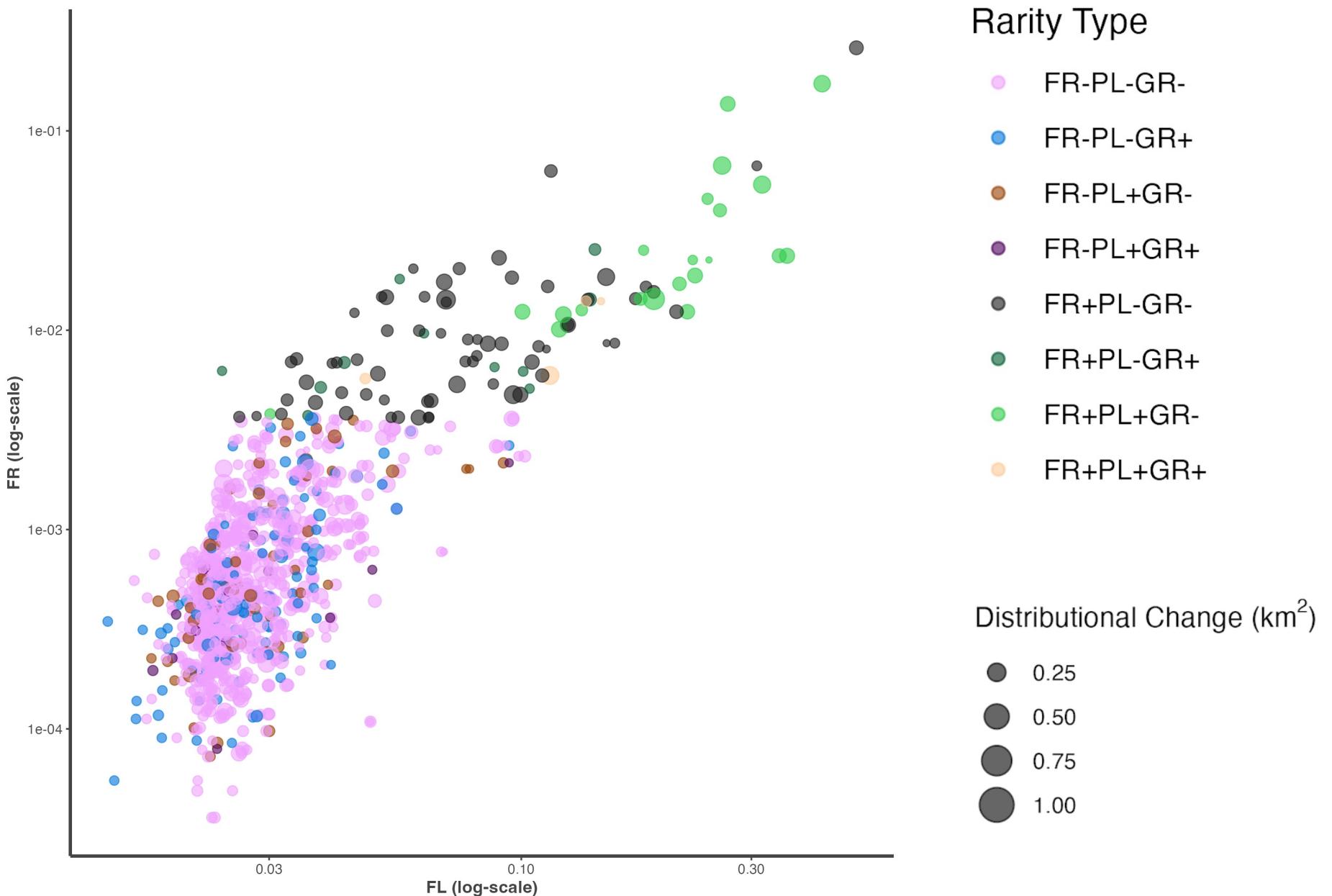
Dimension	AIC Importance
GR	100%
GL	100%
FR	69.8%
FL	45.1%
PR	28.7%
PL	18.5%

Distribution
Change

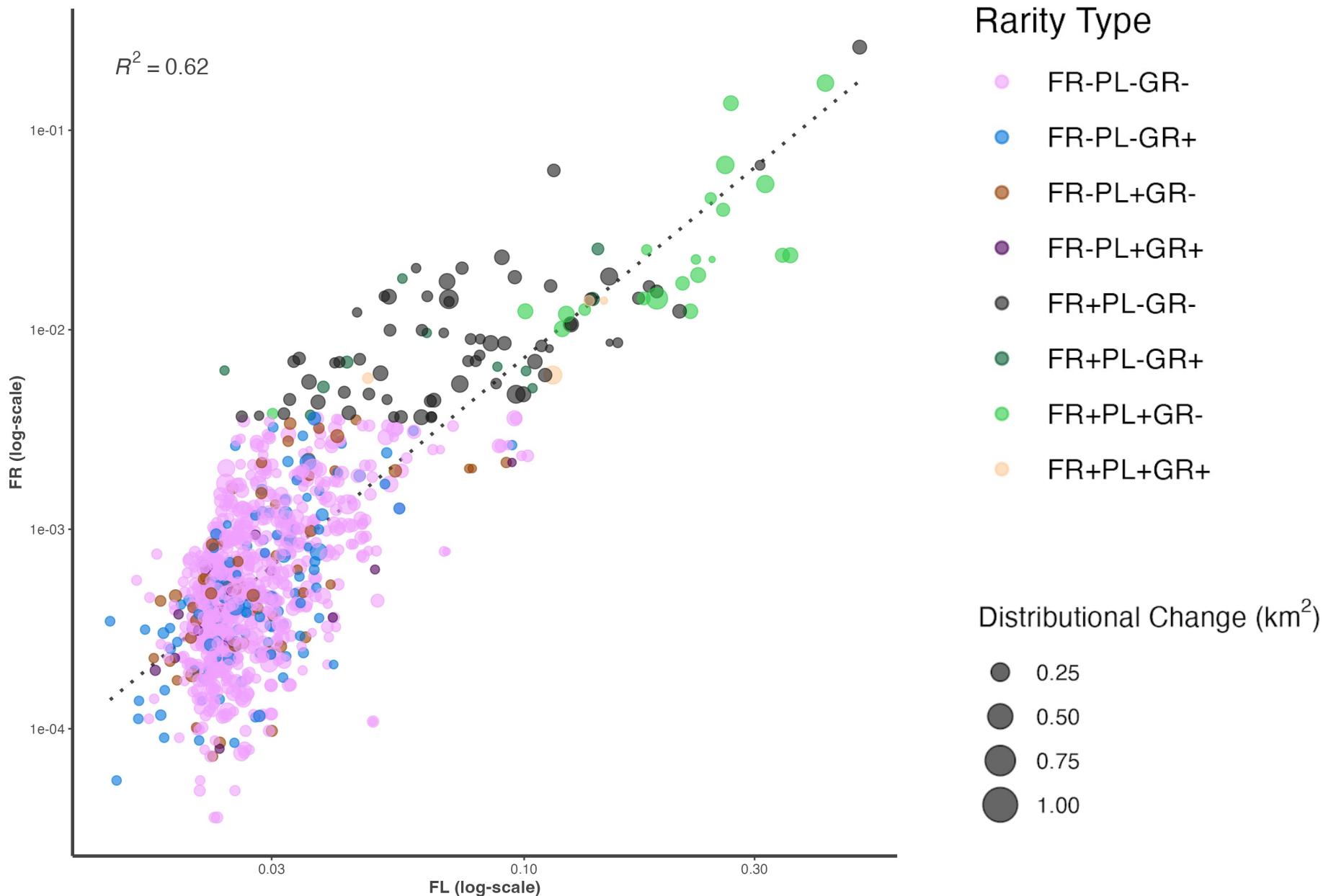
Flowering
Phenology

Dimension	AIC Importance
GL	100%
FR	100%
FL	99.6%
PR	56.3%
PL	23.9%
GR	8.0%

Distribution Change

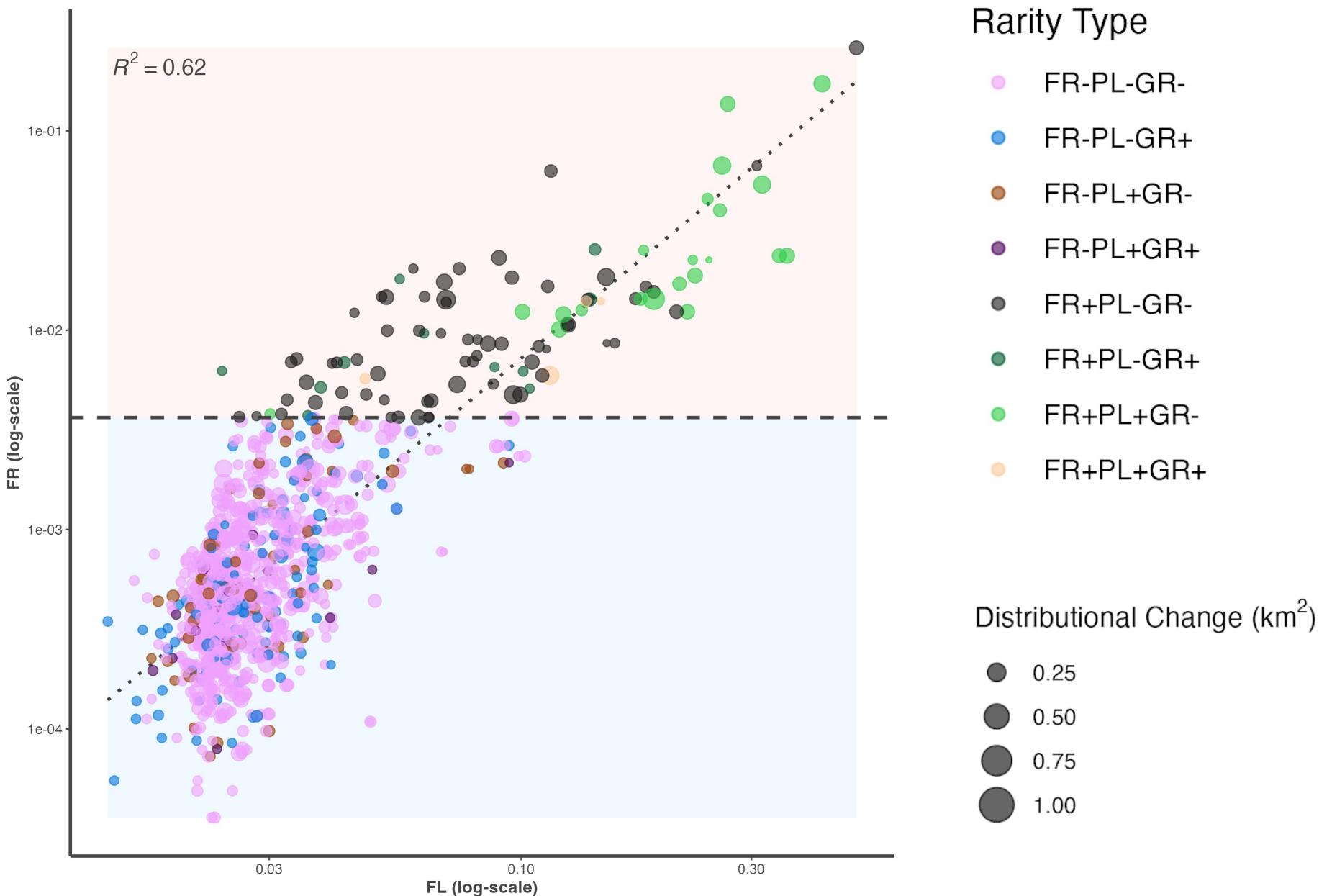


Distribution Change



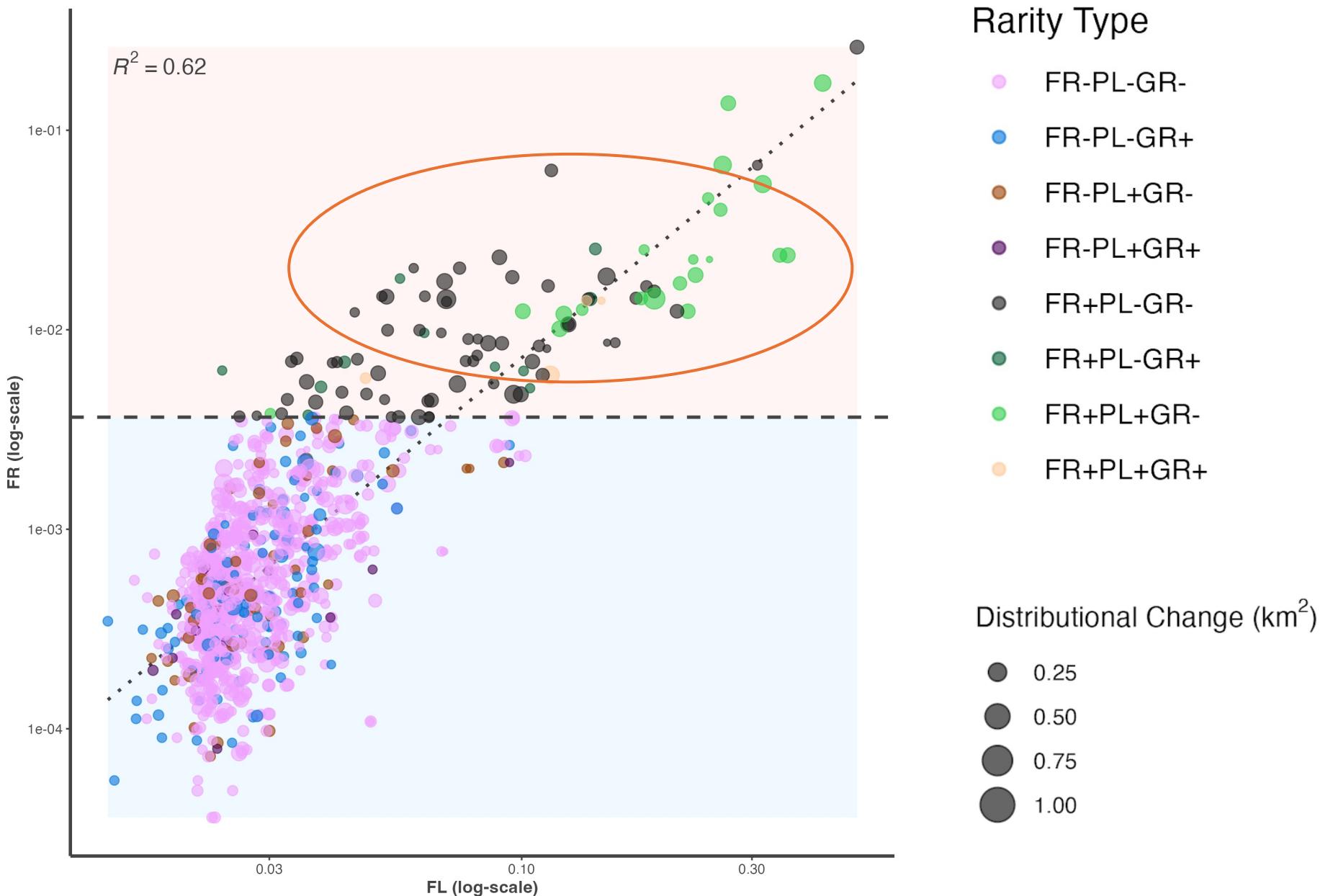
Distribution Change

$G_R F_R P_L$



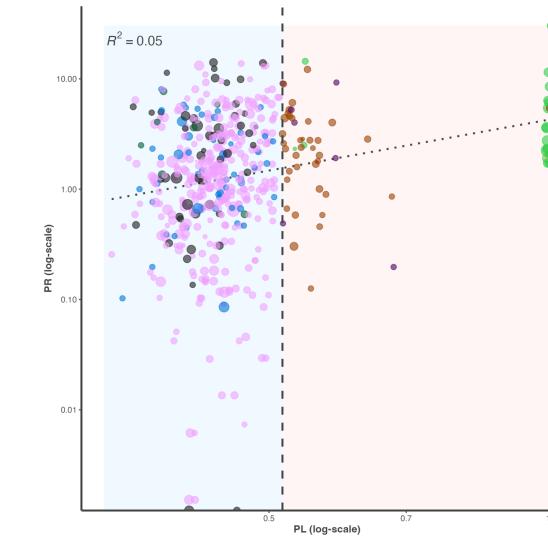
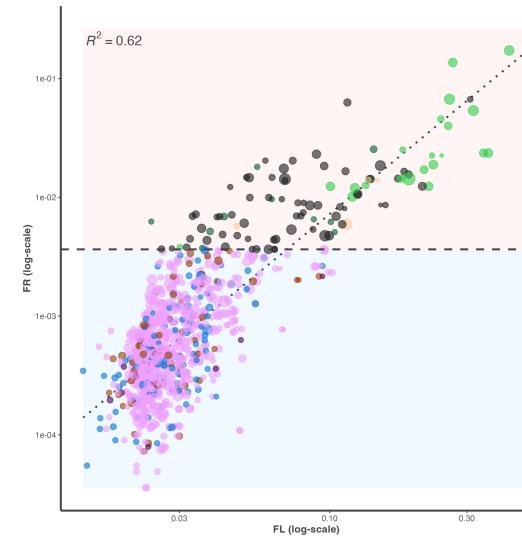
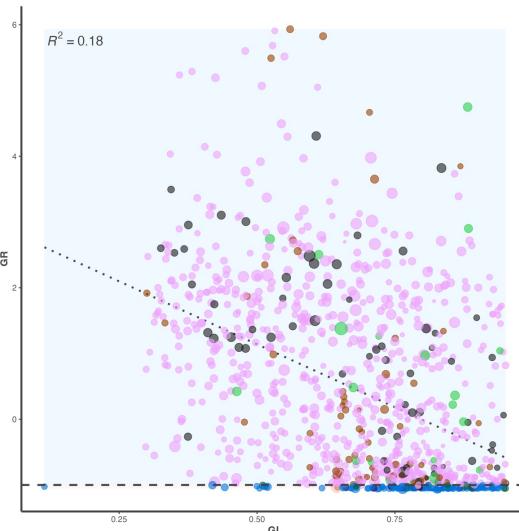
Distribution Change

$G_R F_R P_L$



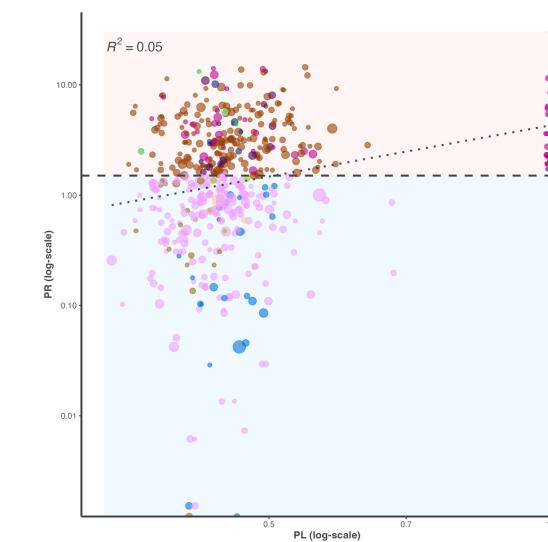
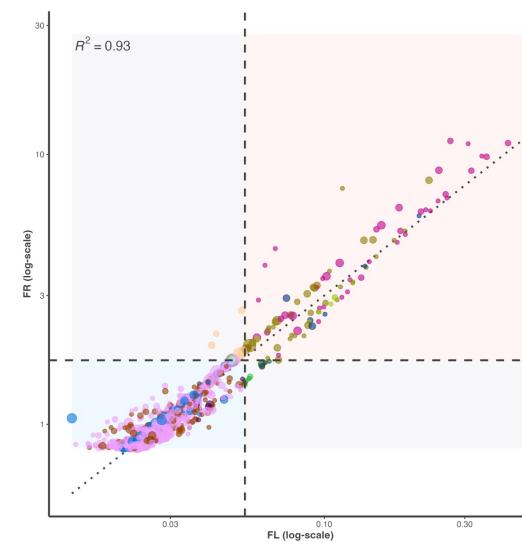
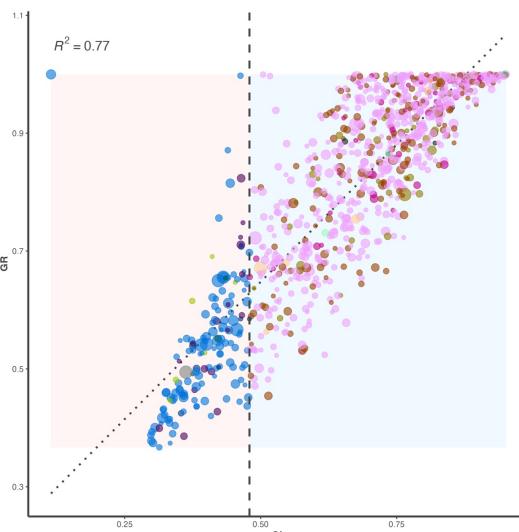
Distribution Change

$G_R F_R P_L$



Flowering Phenology

$G_L F_{RL} P_R$



Rarity Type

- FR-PL-GR-
- FR-PL-GR+
- FR-PL+GR-
- FR-PL+GR+
- FR+PL-GR-
- FR+PL-GR+
- FR+PL+GR-
- FR+PL+GR+

Distributional Change (km^2)

- 0.25
- 0.50
- 0.75
- 1.00

Rarity Type

- FR-FL-PR-GL-
- FR-FL-PR-GL+
- FR-FL-PR+GL-
- FR-FL-PR+GL+
- FR-FL+PR-GL-
- FR-FL+PR-GL+
- FR+FL-PR-GL-
- FR+FL-PR+GL-
- FR+FL+PR-GL-
- FR+FL+PR+GL-
- FR+FL+PR+GL+
- FR+FL+PR+GL+

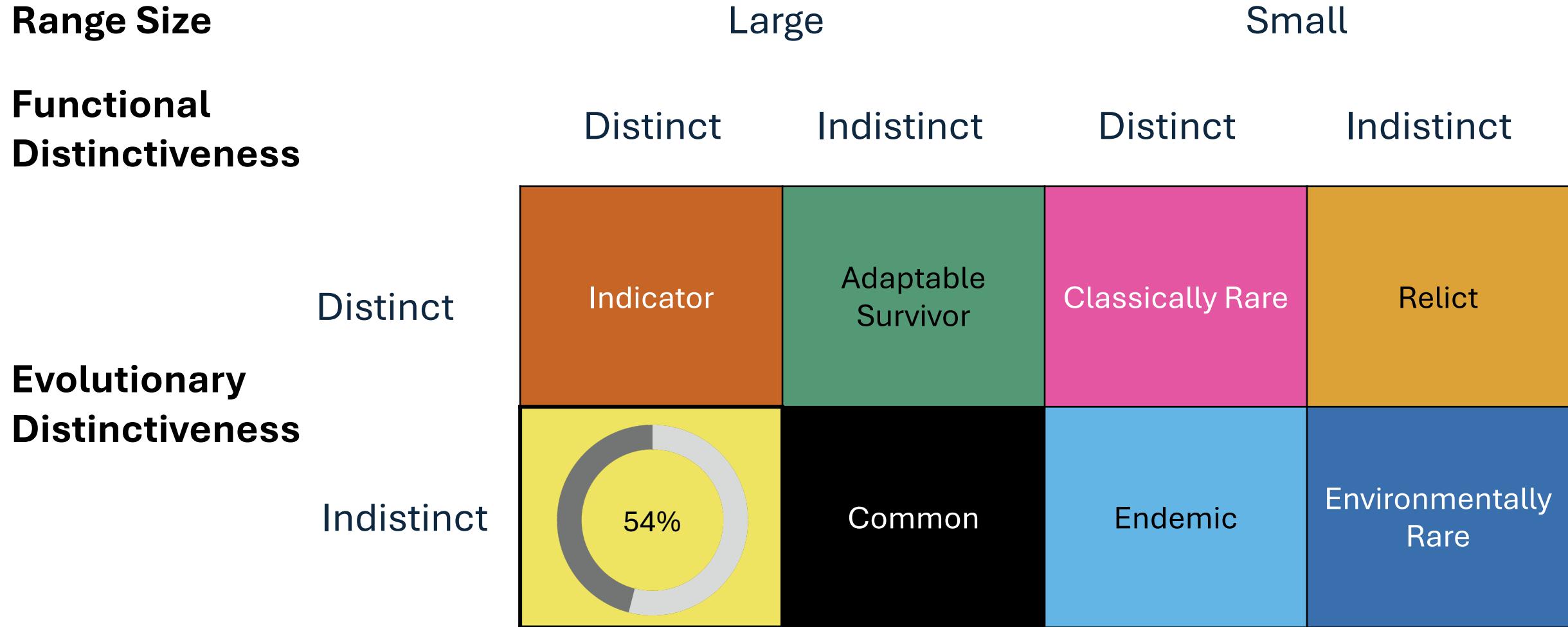
Flowering Window (Months)

- 2.5
- 5.0
- 7.5
- 10.0

EER of Global Angiosperms

Range Size	Large		Small	
Functional Distinctiveness	Distinct	Indistinct	Distinct	Indistinct
Evolutionary Distinctiveness	Distinct	Indicator (0.9%)	Adaptable Survivor (7.4%)	Classically Rare (0.4%)
Evolutionary Distinctiveness	Indistinct	High Invasive Potential (6.3%)	Common (70.4%)	Environmentally Rare (10.9%)

Invasiveness of EER Types



Indicator

Eucalyptus exserta



Queensland Native Seeds
alternativeseeds.com.au

Sun et al. 2016

High Invasive Potential
Pinus radiata



Eugene Zelenko

Adaptable Survivor

Hibbertia riparia



Hammer et al. 2025

Classically Rare

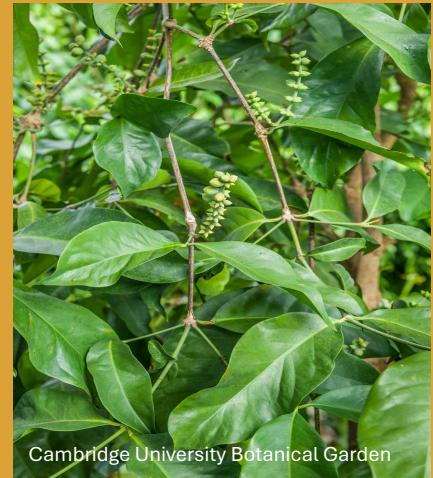
Paulownia kawakamii



Tony Kirkham

Relict

Gnetum gnemon



Cambridge University Botanical Garden

Ickert-Bond et al. 2016

Common
Acer rubrum



Williams & Wardle 2007

Endemic
Flindersia brayleyana



Margaret Donald

Boland 2006

Environmentally Rare
Aporosa frutescens



Lam et al. 2022

1

Can we use rarity as a
tool to understand
ecological processes?
How?

2

Are rarity types robust
and ecologically
relevant?

1

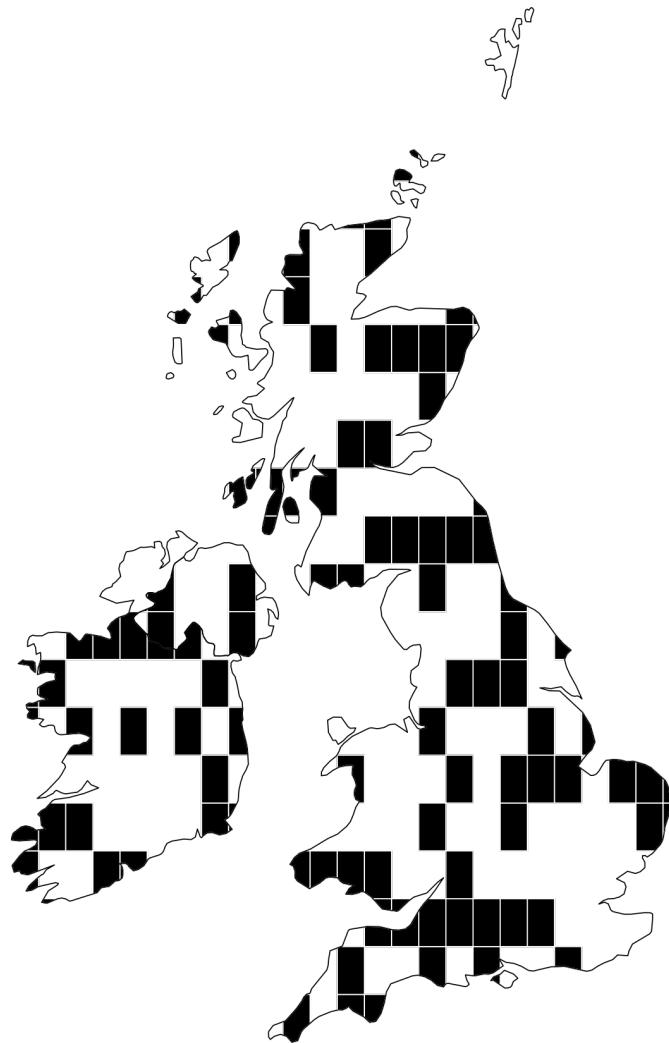
Can we use rarity as a
tool to understand
ecological processes?

How?

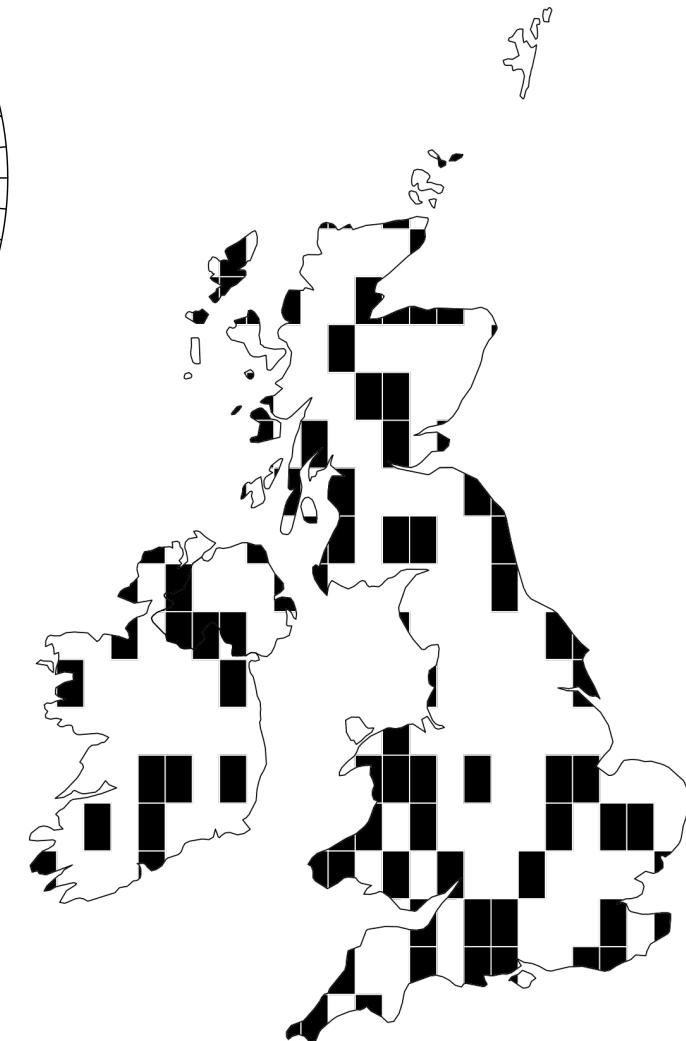
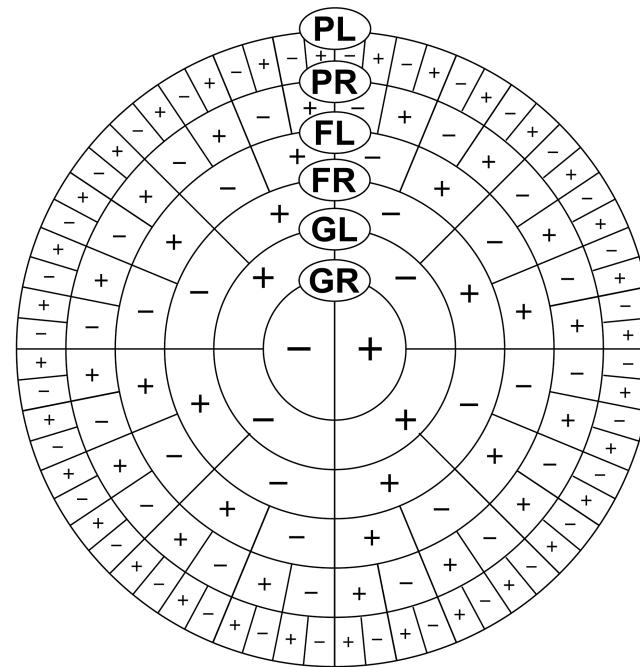
**Yes... with some
caveats**

2

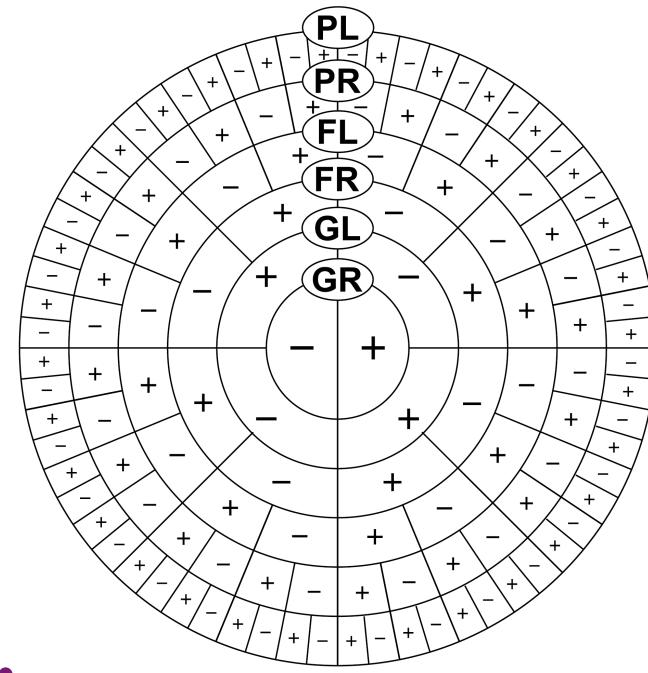
Are rarity types robust
and ecologically
relevant?



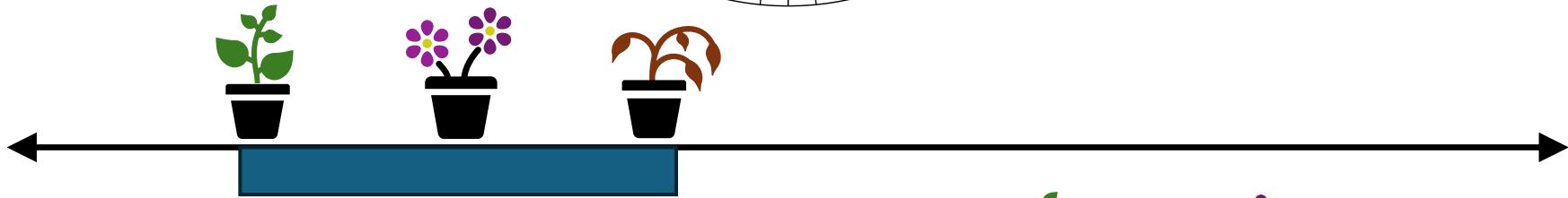
1987



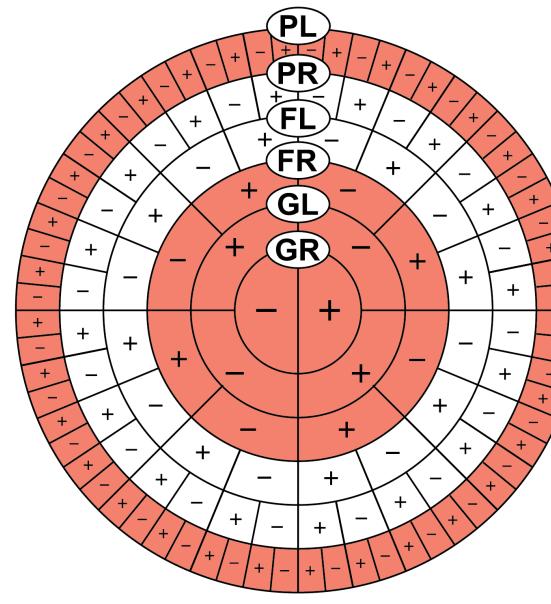
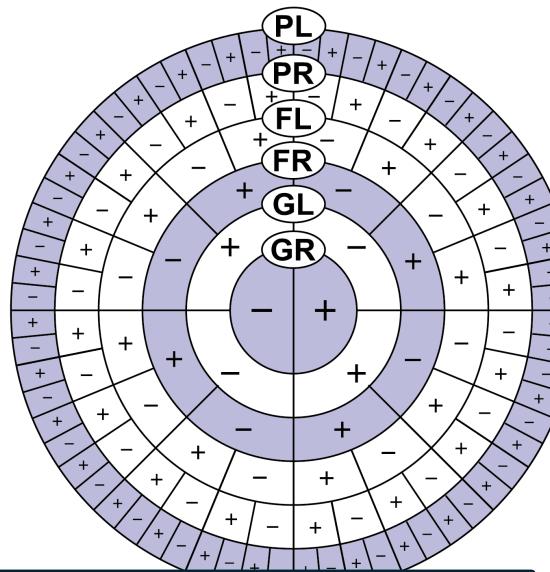
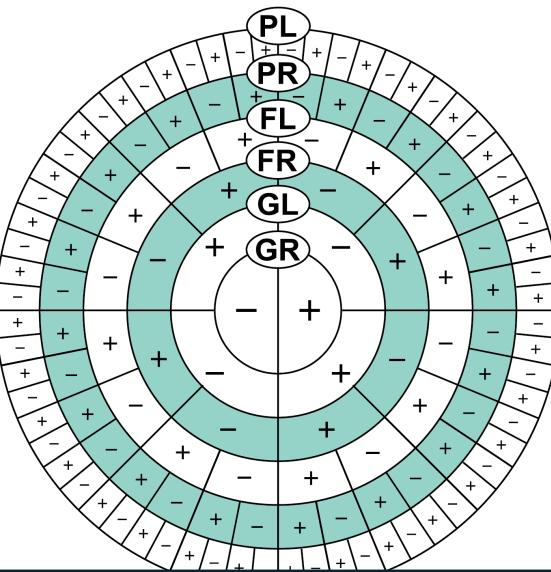
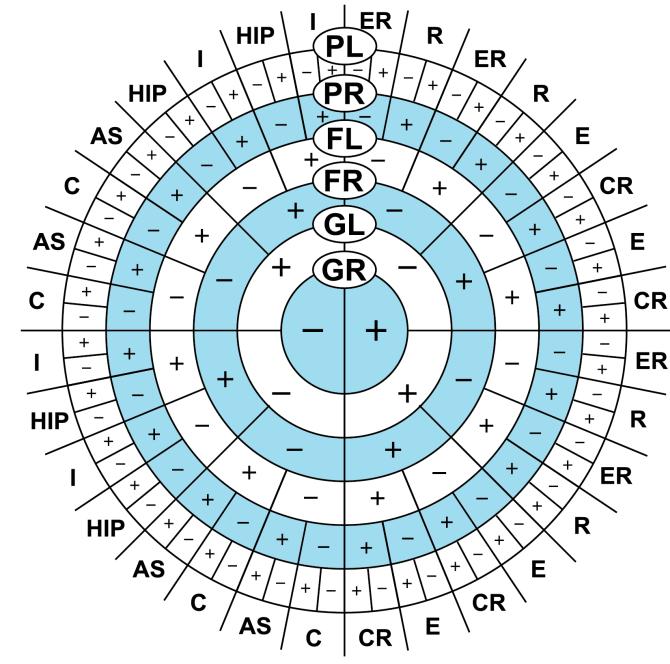
2019



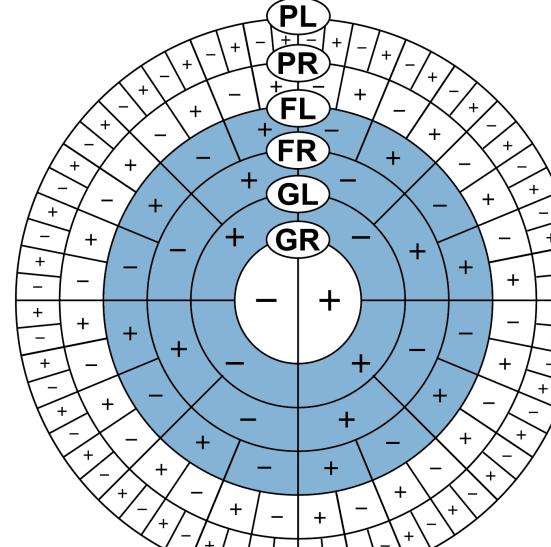
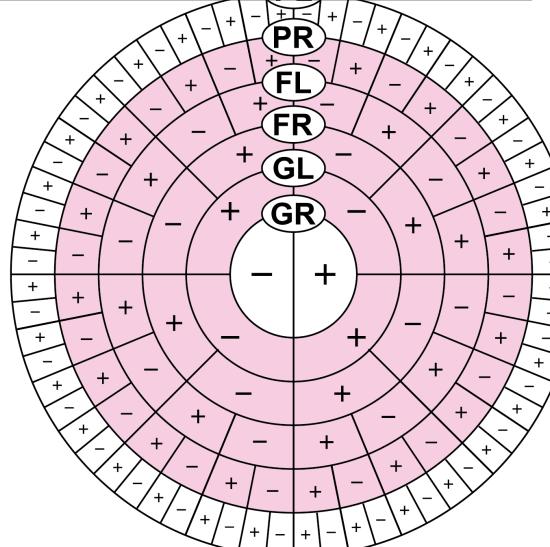
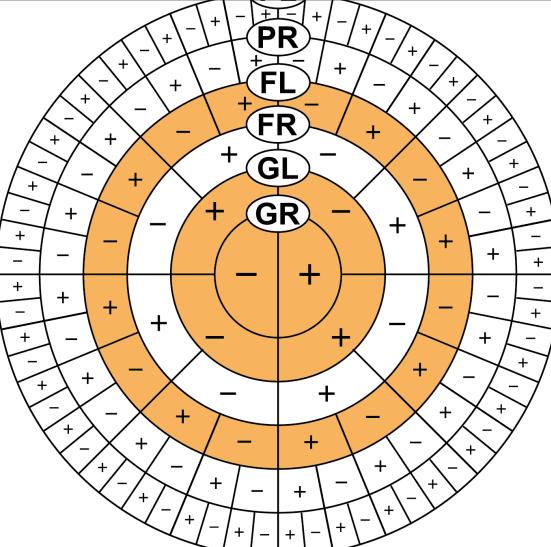
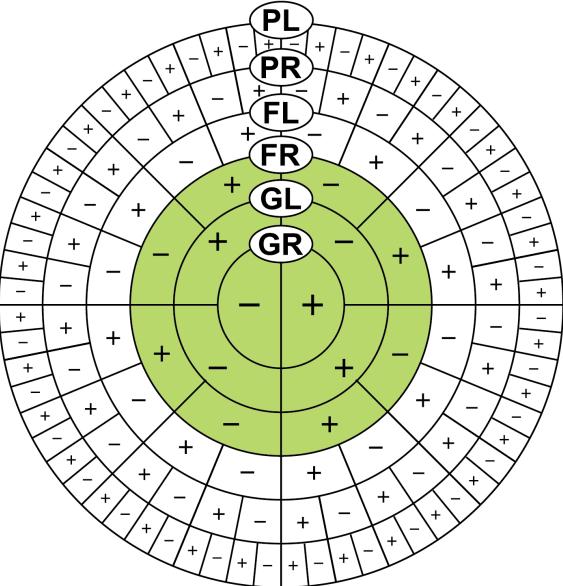
Spring

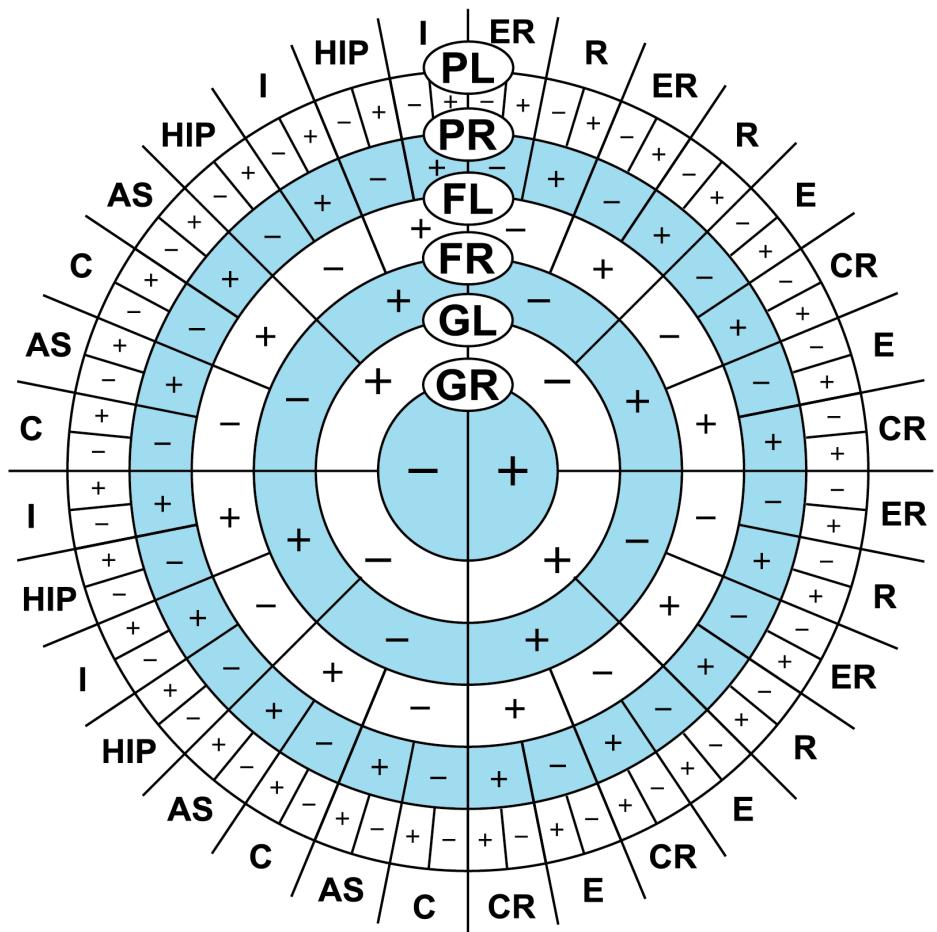


Fall



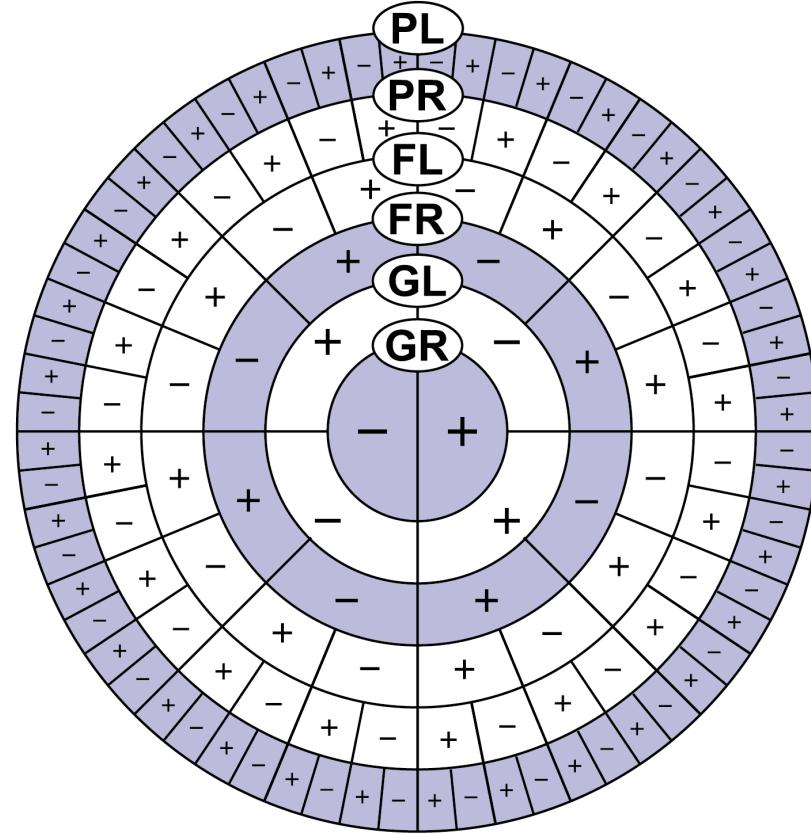
Novel Restrictions



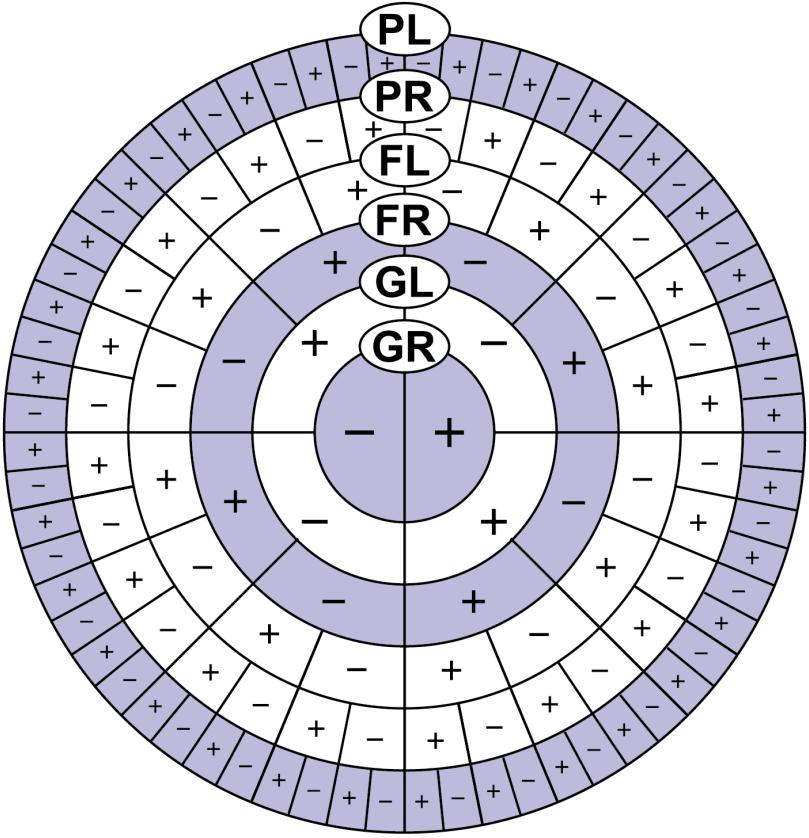


Eco-Evolutionary Rarity

VS

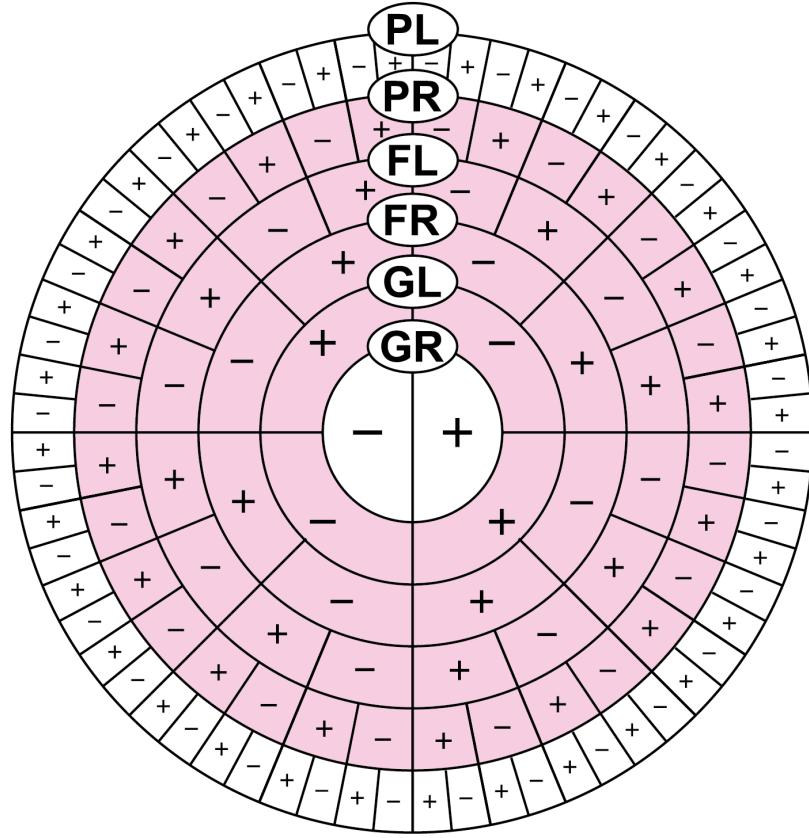


$G_R F_R P_L$

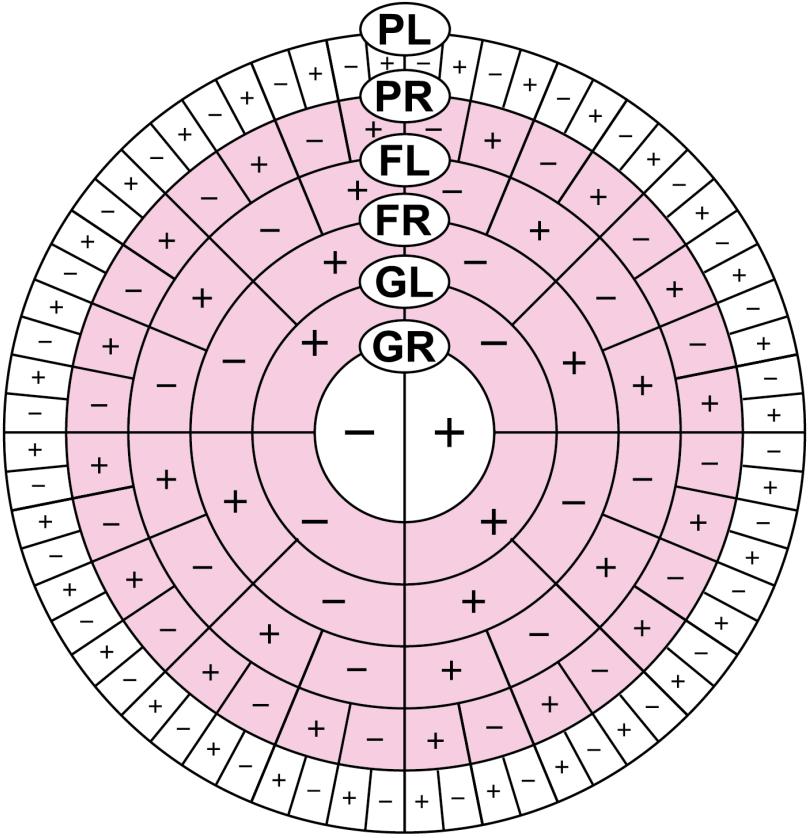


$G_R F_R P_L$

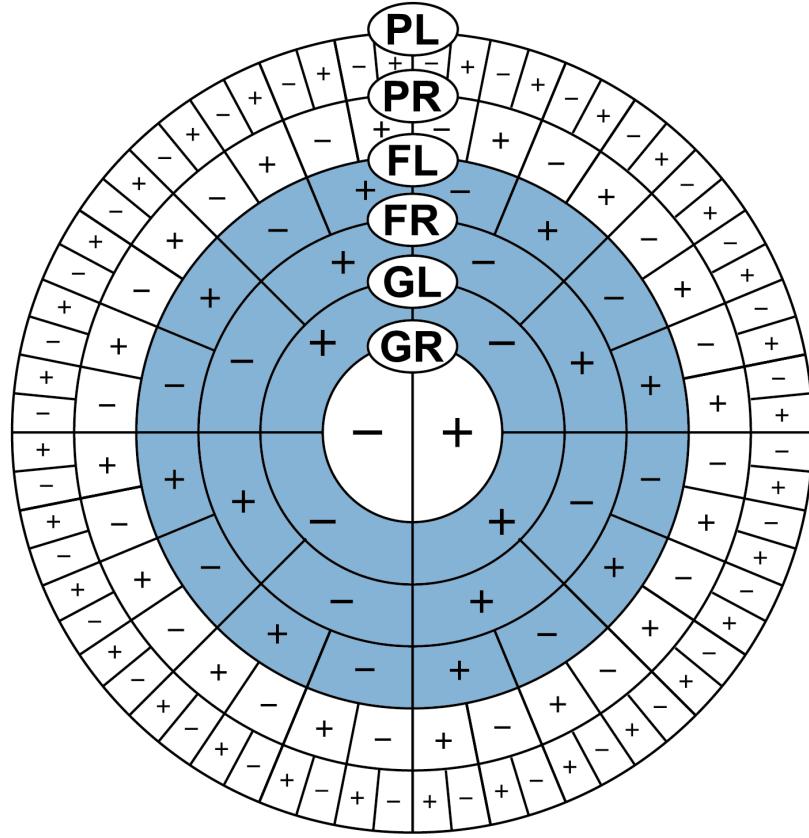
VS



$G_{RL} F_{RL} P_R$

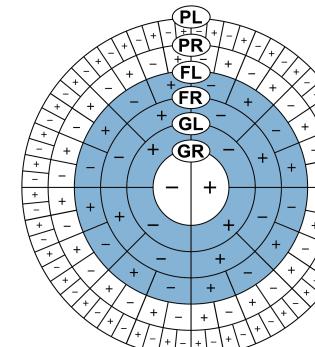
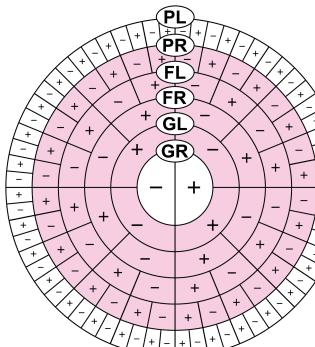
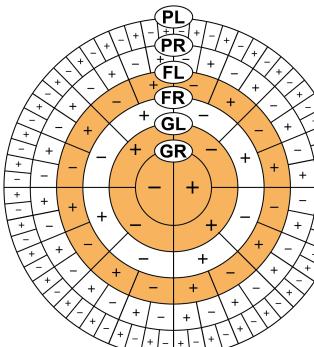
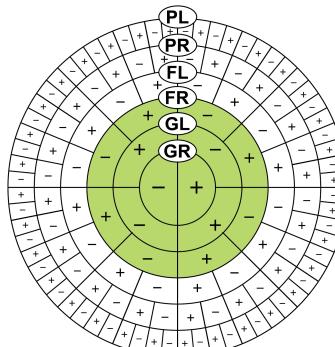
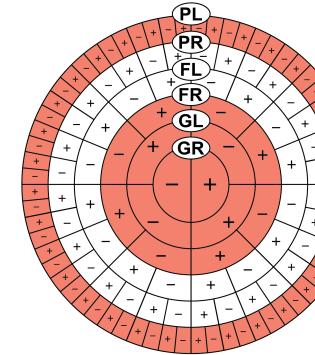
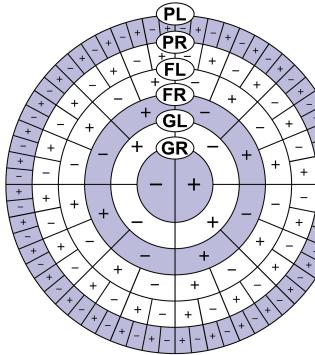
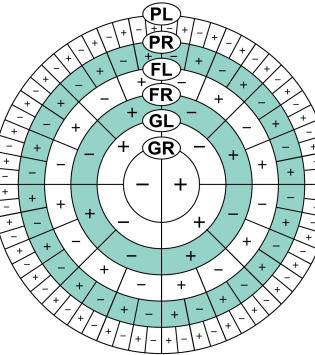
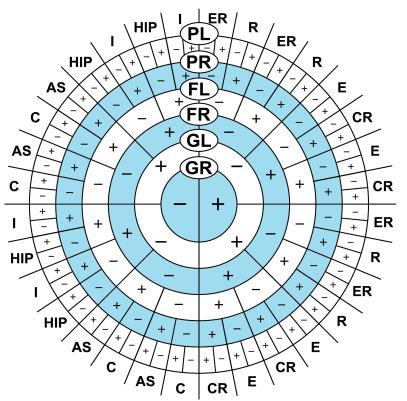
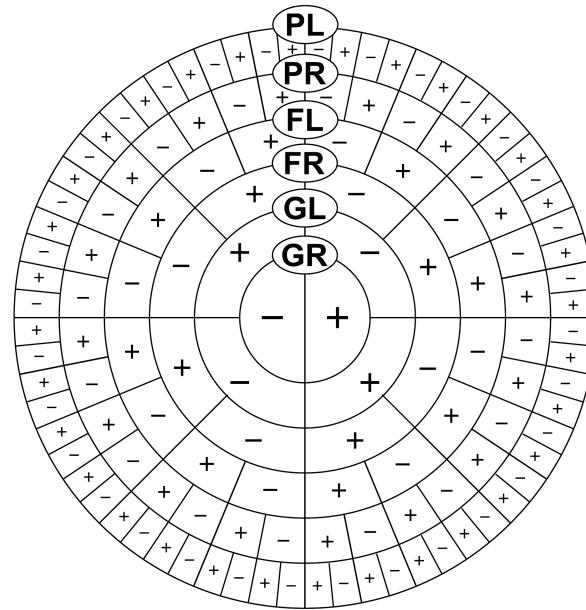


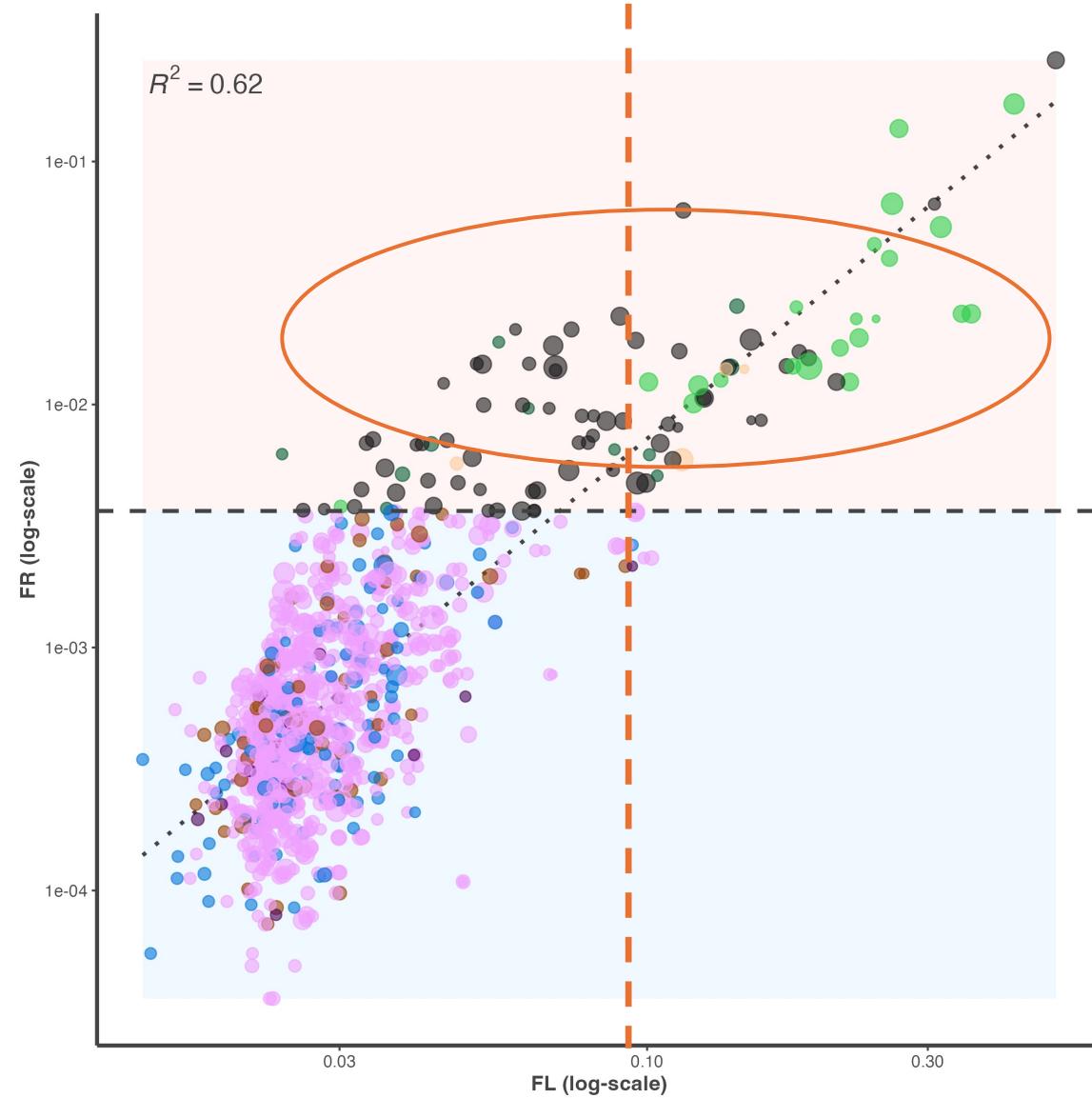
VS



$G_{RL}F_{RL}P_R$

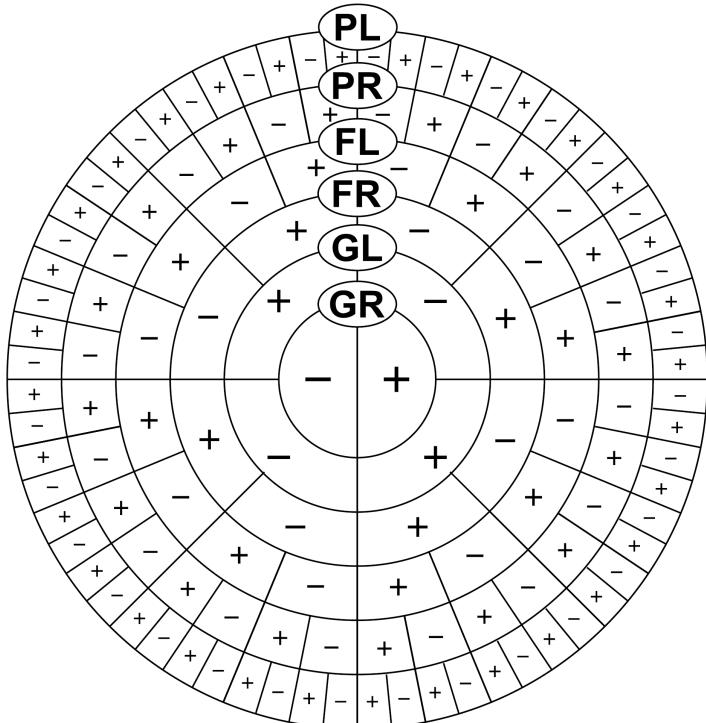
G_LF_{RL}



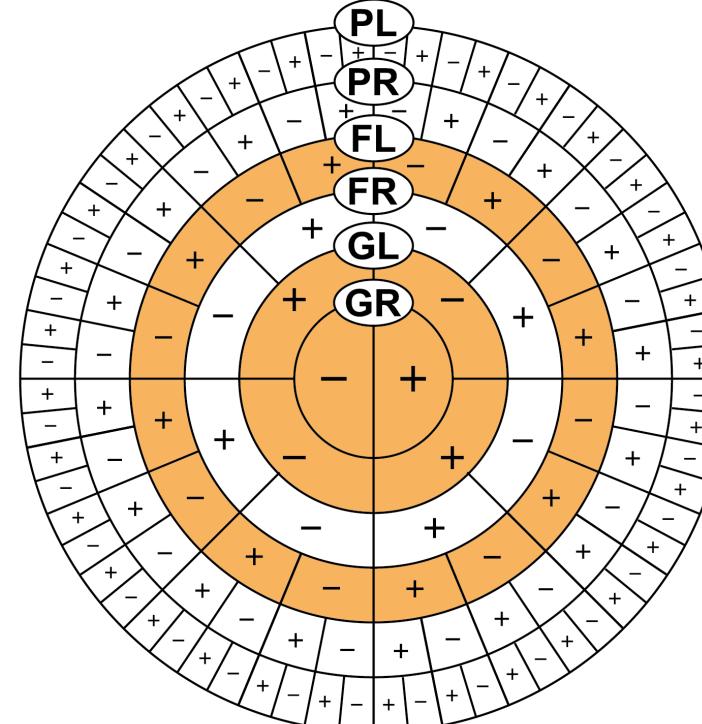
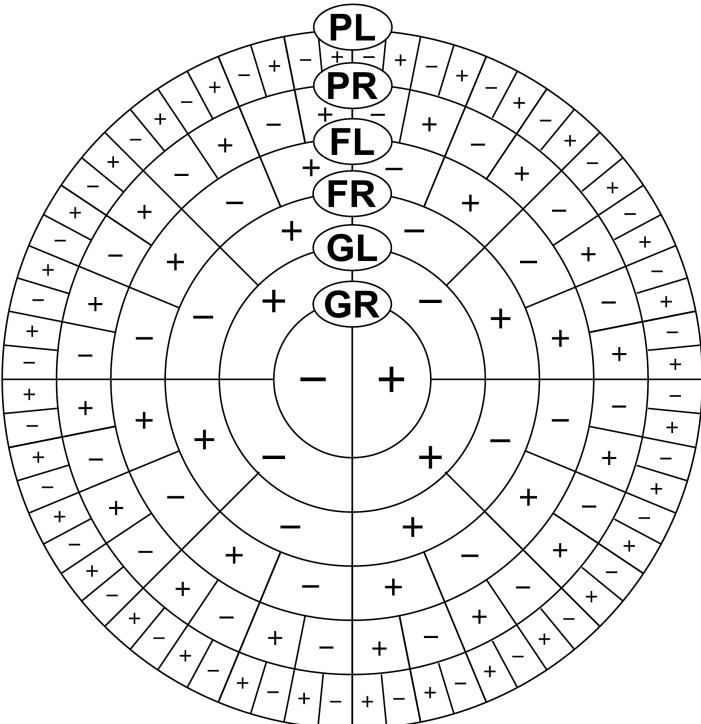


When and if rarity is best defined by its
comprehensiveness or by its utility in
explaining ecology?

When and if rarity is best defined by its comprehensiveness or by its utility in explaining ecology?



When and if rarity is best defined by its comprehensiveness or by its utility in explaining ecology?



63 Restrictions

GRFRLPRL
GRLPRL EER GRFRLPRL
GRFRLPL GRLPL
GRFRL FRL
GRFLR GRFLPL FLPL GRLFL FRL
GRFL FRPR FRLPRL GRLFRPRL GLFRLPL
GRFRPL GLFL GRLFR PRL EDGE
GRLFLPL GLFRPR GLFLPR GLFLPL GRLFRLPL
GRL GRLPRL PR GRFLPR GRPRL
FLPRL GLPR FL GDR GLFRL FRLPR FRLPL
GLFLPRL GL FLPR GLFRLPRL
GPL GLPRL FRPRL
GR Functional Rarity GLFRLPR
FR FRPL GRLFRPR GRFRPRL
GRLFRPRL GRFR GRLFRPR PL GRFLPRL
GLFRPRL GRLFLPR

Limiting Abundance Data

GRFRLPR
GRFLPRL EER GRFRLPRL
GRFRLPL GRLFRLPR GRLPL
GRFRL GRFLPL FRL
GRFL FRPR FLPL GRLFL GRLFLPRL
GRFRPL GLFL FRLPRL GRLFRPRL GLFRLPL
GRLFLPL GLFRPR GRLFLPR PRL EDGE
GRLFR PR GLFLPL GRLFRLPL
GRL GRLFRPRL GLFLPR PR GRFLPR GRPRL
FLPRL GLPR FL GDR GLFRL FRLPR FRLPL
GLFLPRL GL FLPR FRPRL GLFRLPRL
GPL GLPRL Functional Rarity GLFRLPRL
GR FR FRPL GRLFRPR GRFRPRL
GRLFRPRL GRFR GRPL GRFLPRL
GLFRPRL GRLFLPR

Defining Rarity

GRFRPL **EER** **GRFRLPRL**
GRFRLPL GRLPRL GRLFLPR
GRFRL GRFLPL FRL
GRFL FRPR GRLFLPL
GRLFLPL GLFL FRLPRL GLFRLPL
GRLFR GRLFRPRL EDGE
GLFLPL GLFRPR GRLFRPL GRLFRLPL
GRL GRLFRPL GLFLPR PR GRFLPR
FLPRL GLPR FL GDR GLFRL FRLPR FRLPL
GLFLPRL GL FLPR GLFRLPRL
GPL GPLR FRPRL GLFRLPRL
GR FR FRPL Functional Rarity GLFRLPR
GRLFRPL GRFR GRLFRPR GRFRPRL
GLFRPRL PL GRFLPRL
GRLFLPR

Explaining Range Dynamics

GRFRPL

EER

GRFRLPRL

GRLPL

FRL

GRLFLPRL

GLFRLPL

EDGE

GRLFRLPL

PR

GRFLPRL

GRPRL

FRLPL

GLFRLPRL

GLFRLPR

FRPRL

Functional Rarity

GLFLPRL

GL

FLPR

GL

FLPR

GPL

GLPRL

FRPL

GR

FR

GRFR

PL

GRFLPRL

GRLFRPRL

GLFRPRL

GRLFLPRL

GRFLPRL

1

Can we use rarity as a
tool to understand
biological processes?

How?

**Yes... with some
caveats**

2

Are rarity types robust
and ecologically
relevant?

1

Can we use rarity as a
tool to understand
biological processes?

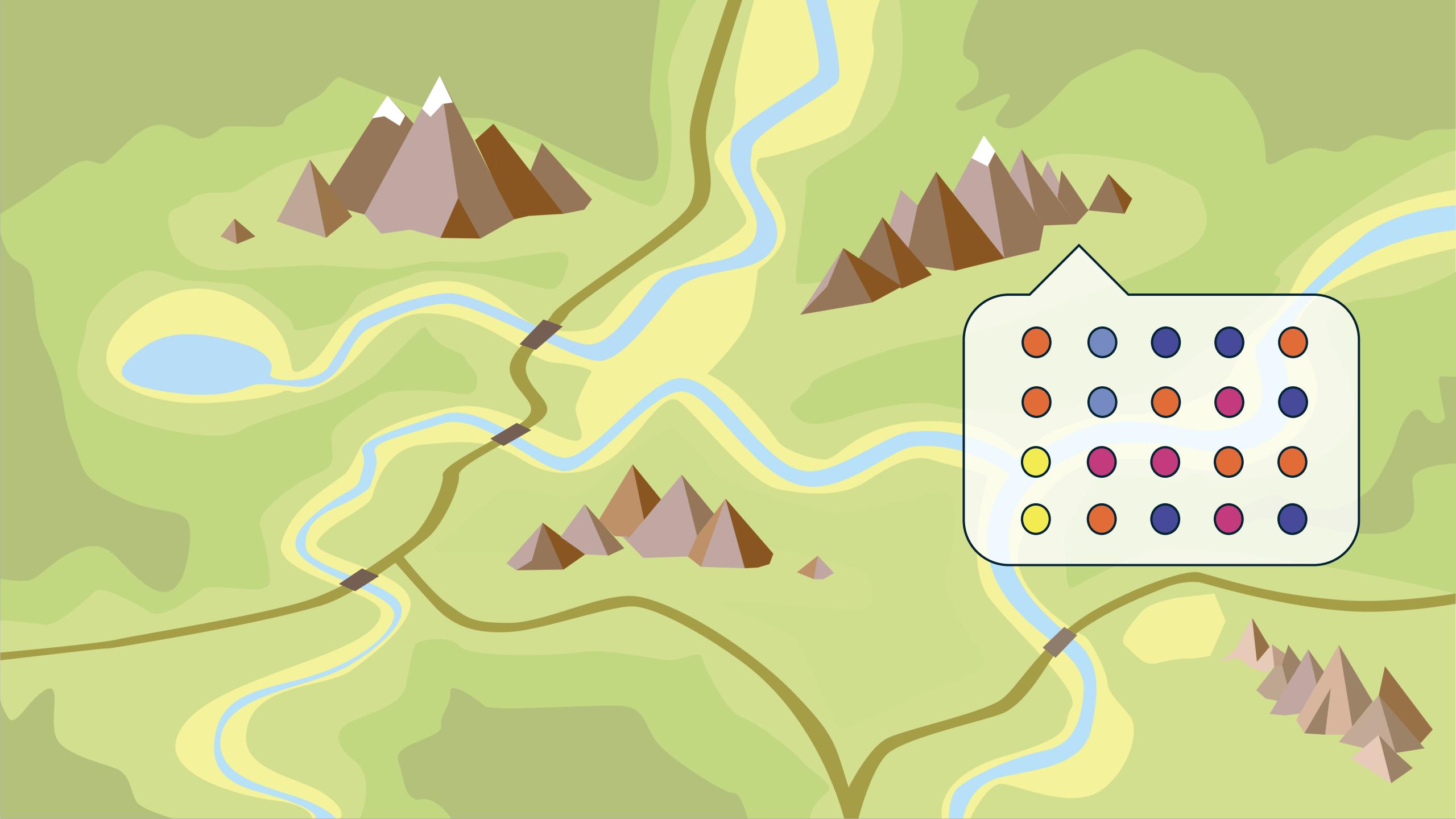
How?

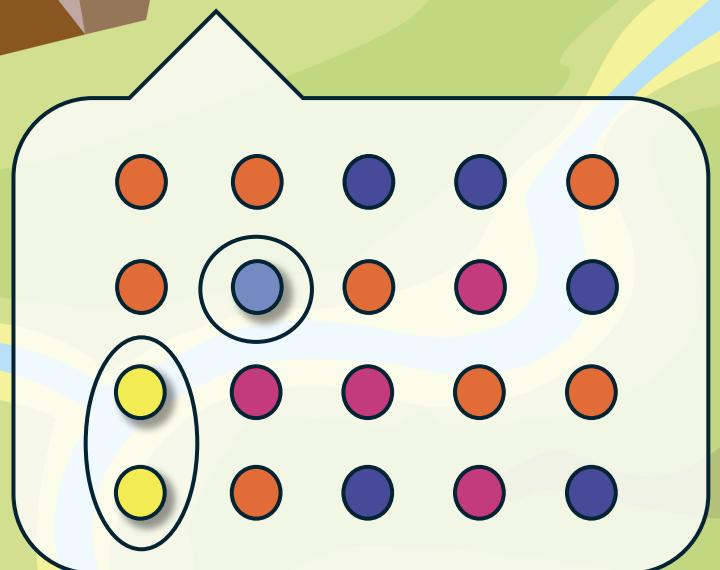
Yes... with some
caveats

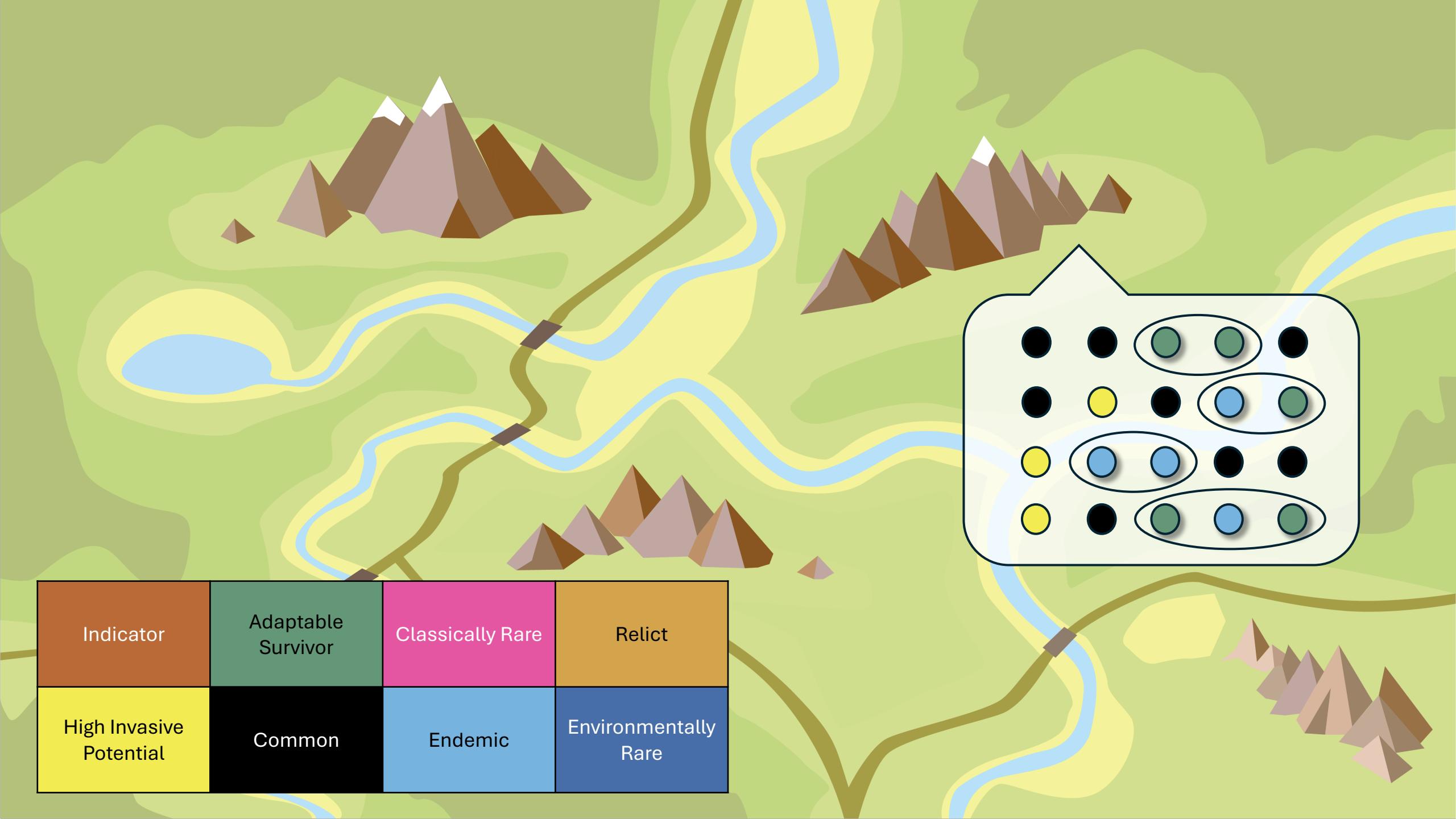
2

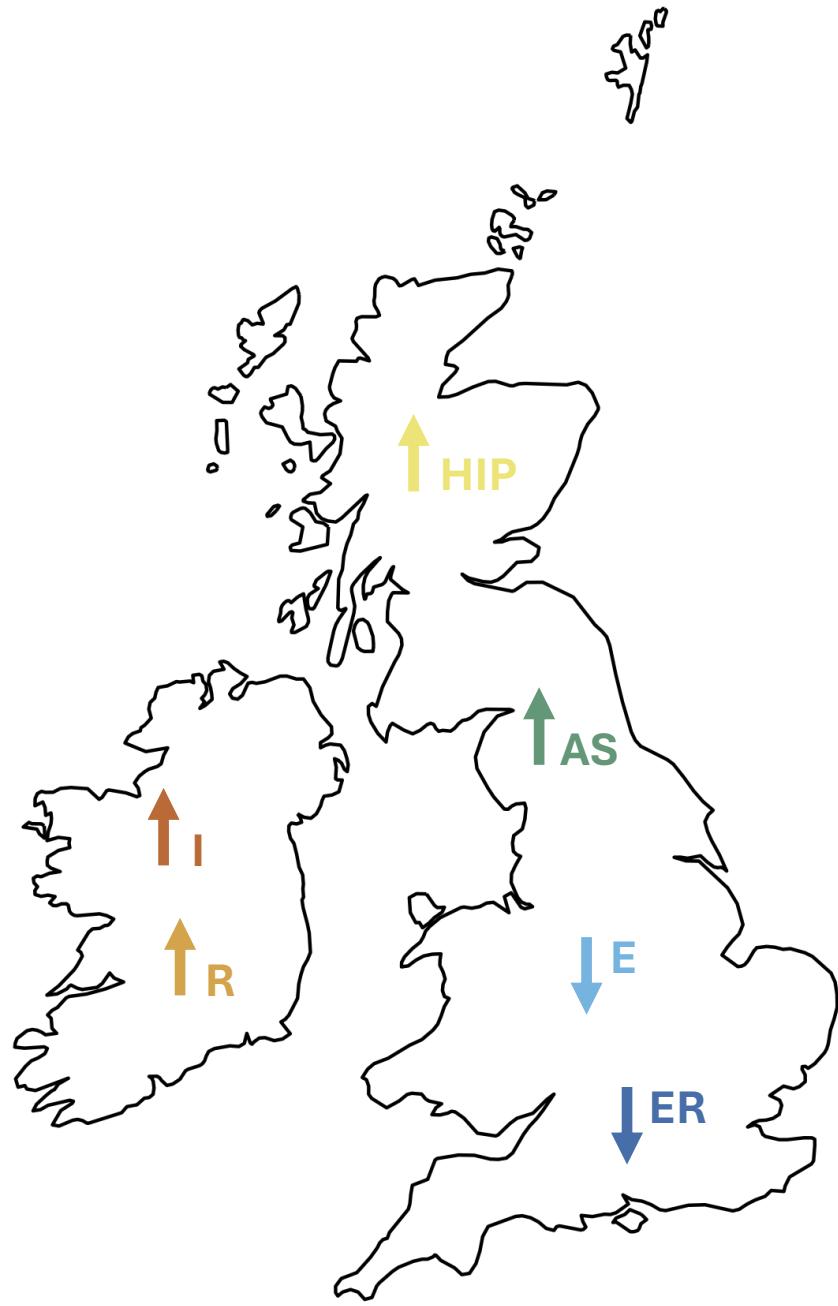
Are rarity types robust
and ecologically
relevant?

Probably... with large
impacts for
conservation

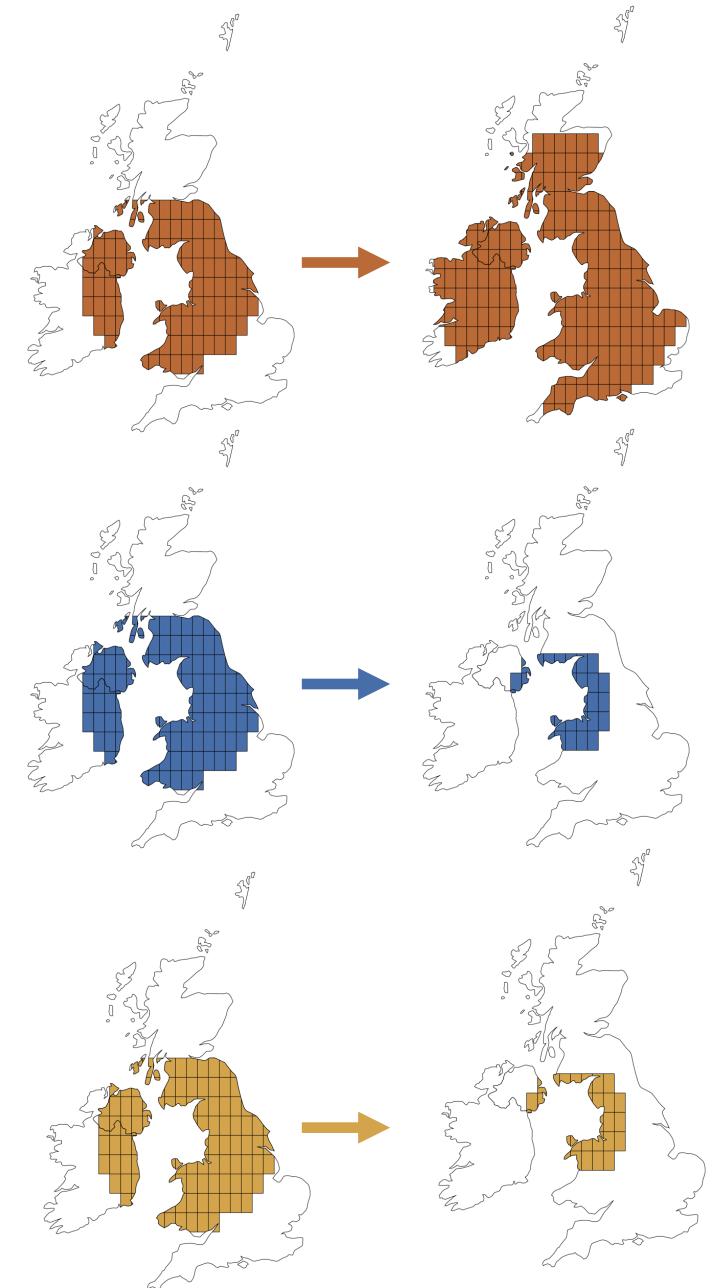








ENDEMIC
RELICT *In-situ
ENVIRONMENTALLY
RARE *Ex-situ



The **Global Model of Discretized Rarity** and its complementary restrictions overcome data limitations and provide more than 50 novel definitions of rarity and tools to understand ecological processes, grounded in a unified conceptual framework.

Future Directions



- Identify a suite of complementary restrictions that capture a wide range of ecological processes.
- Uncover the abiotic and biotic gradients that have shaped diversity in rarity within these restrictions through evolutionary time.
- Link rarity to ecosystem function.

Predict how species, communities, and ecosystems respond to climate change using **biodiversity-rarity-ecosystem function (BREF)** relationships.

Thank you!

Beaulieu
Lab

O'Meara
Lab

Bailey
Lab



anytko@vols.utk.edu