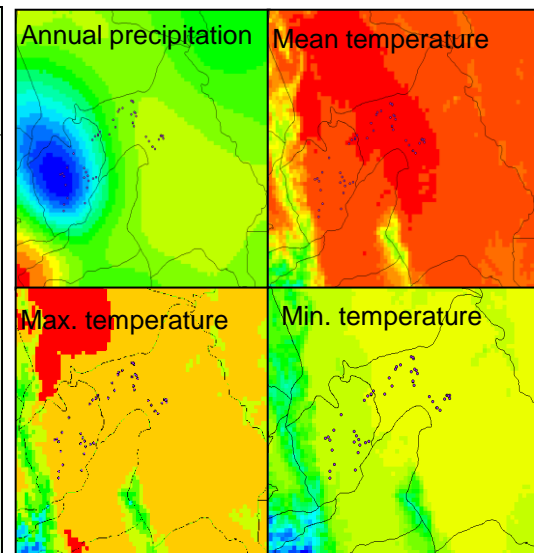
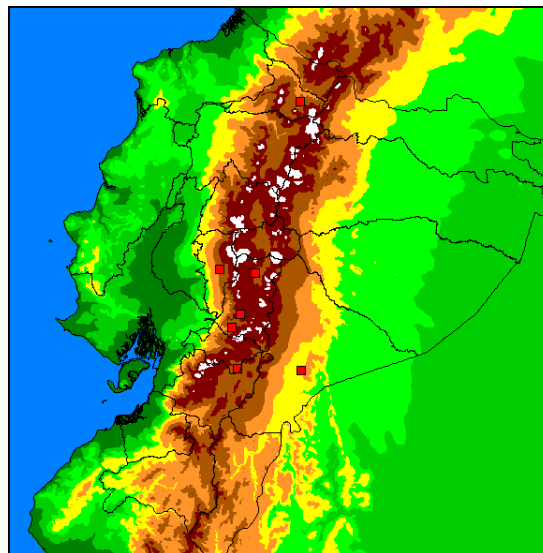
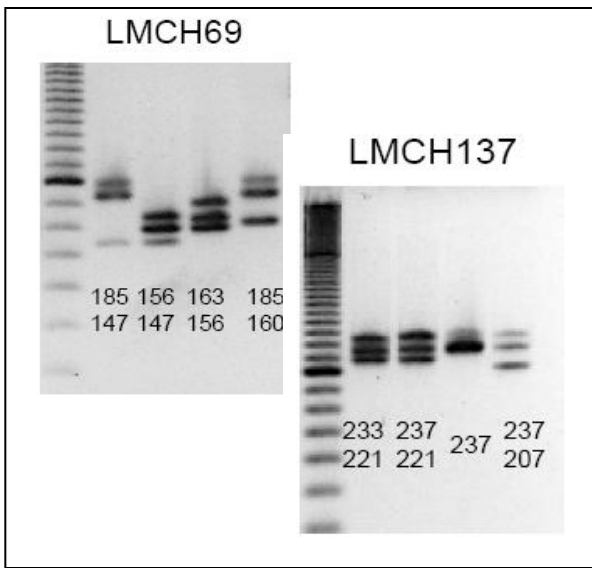
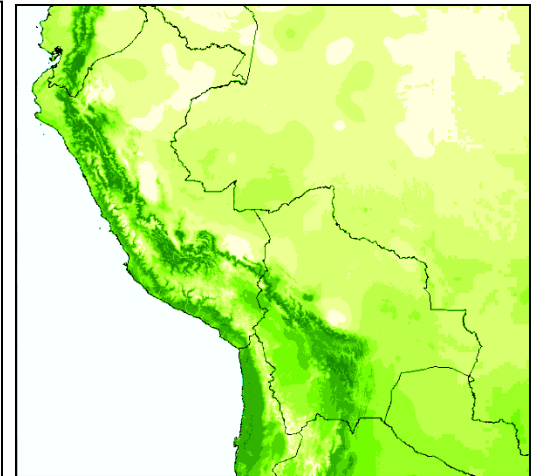
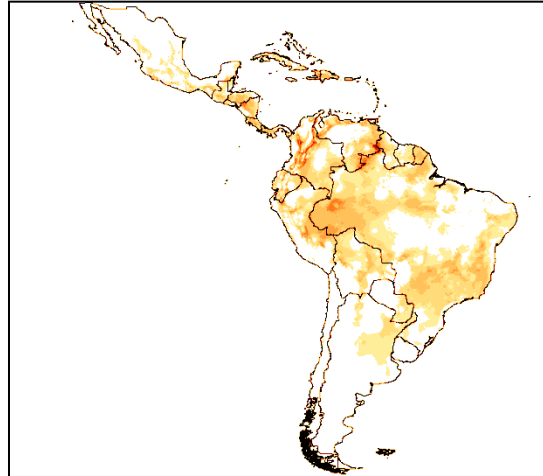
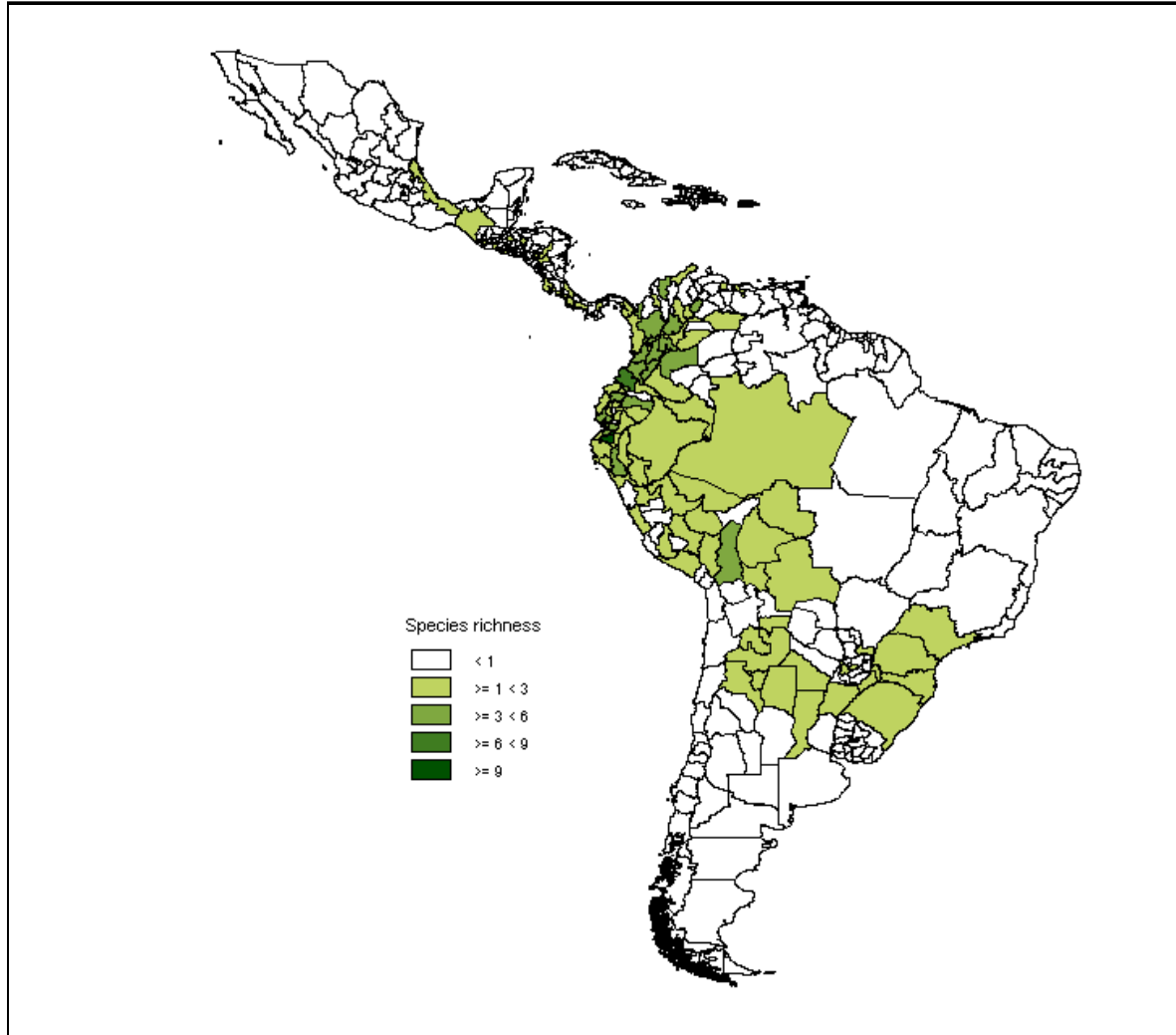


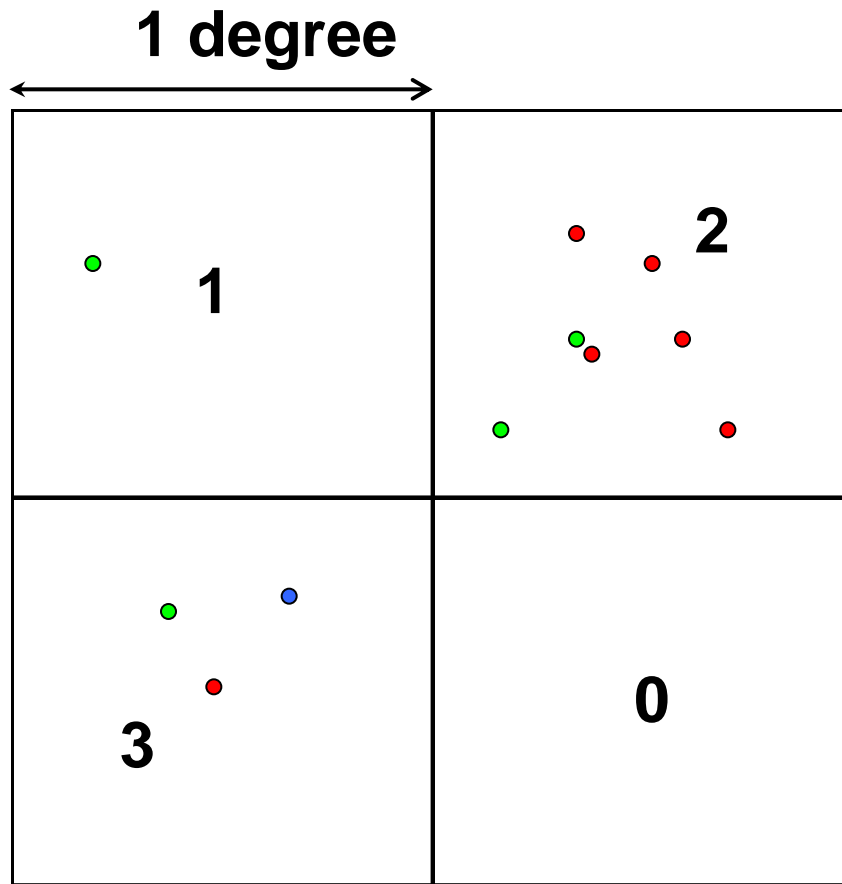
Introduction to diversity analysis and its implications

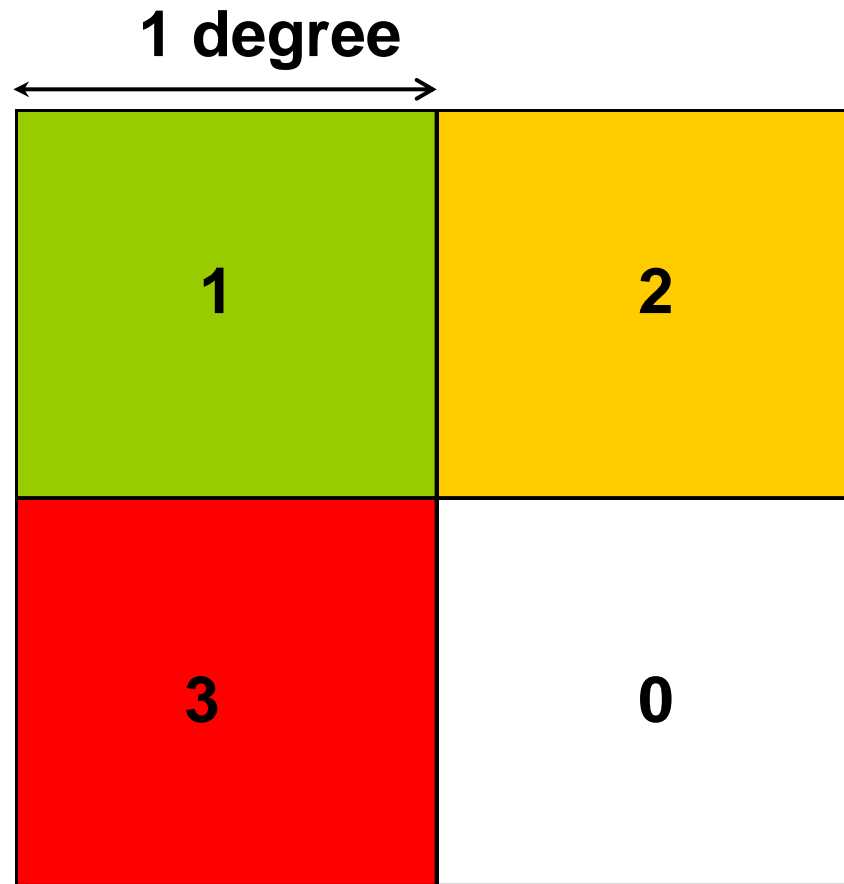


Diversity question...

Where can the highest *Vasconcellea* diversity be found?

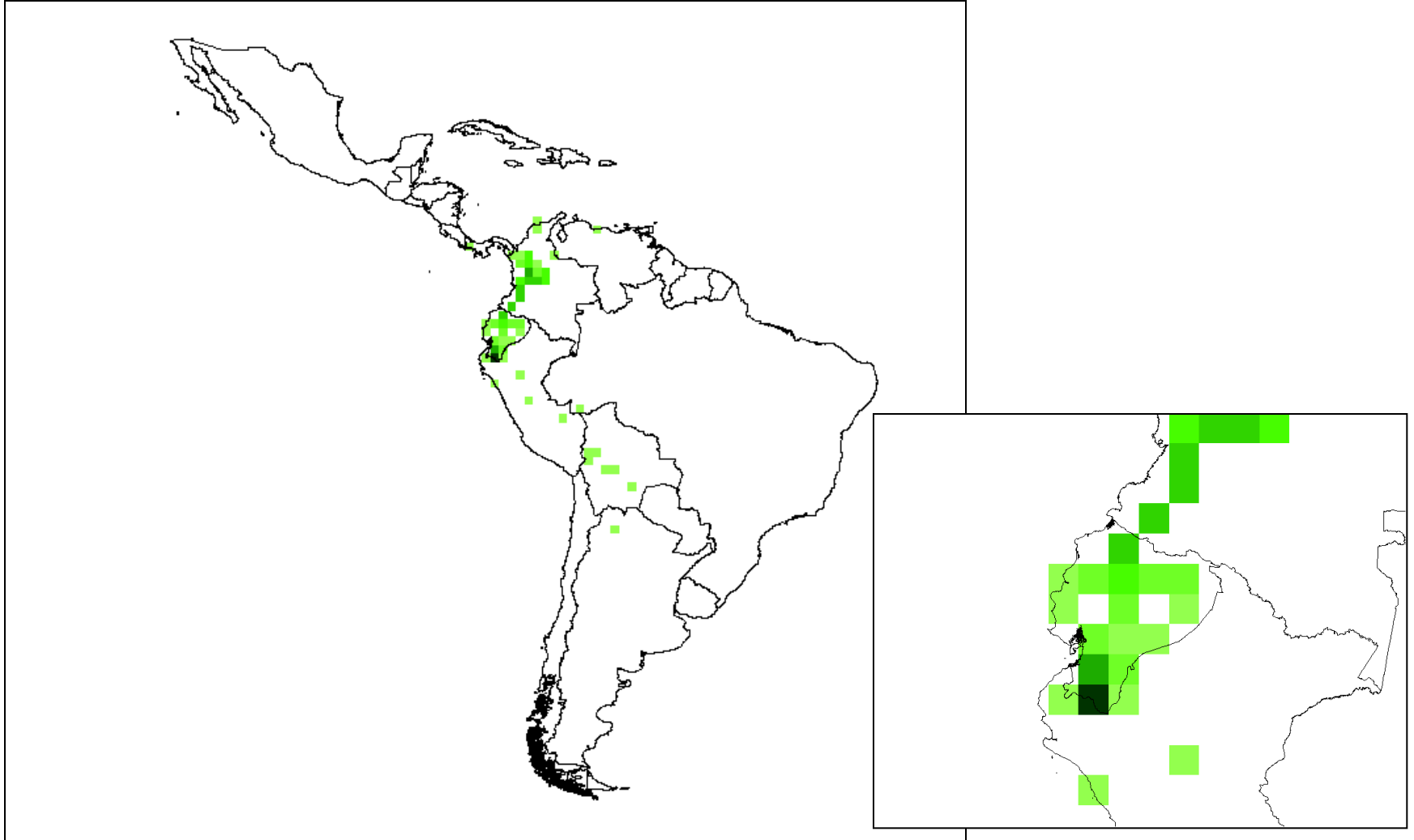


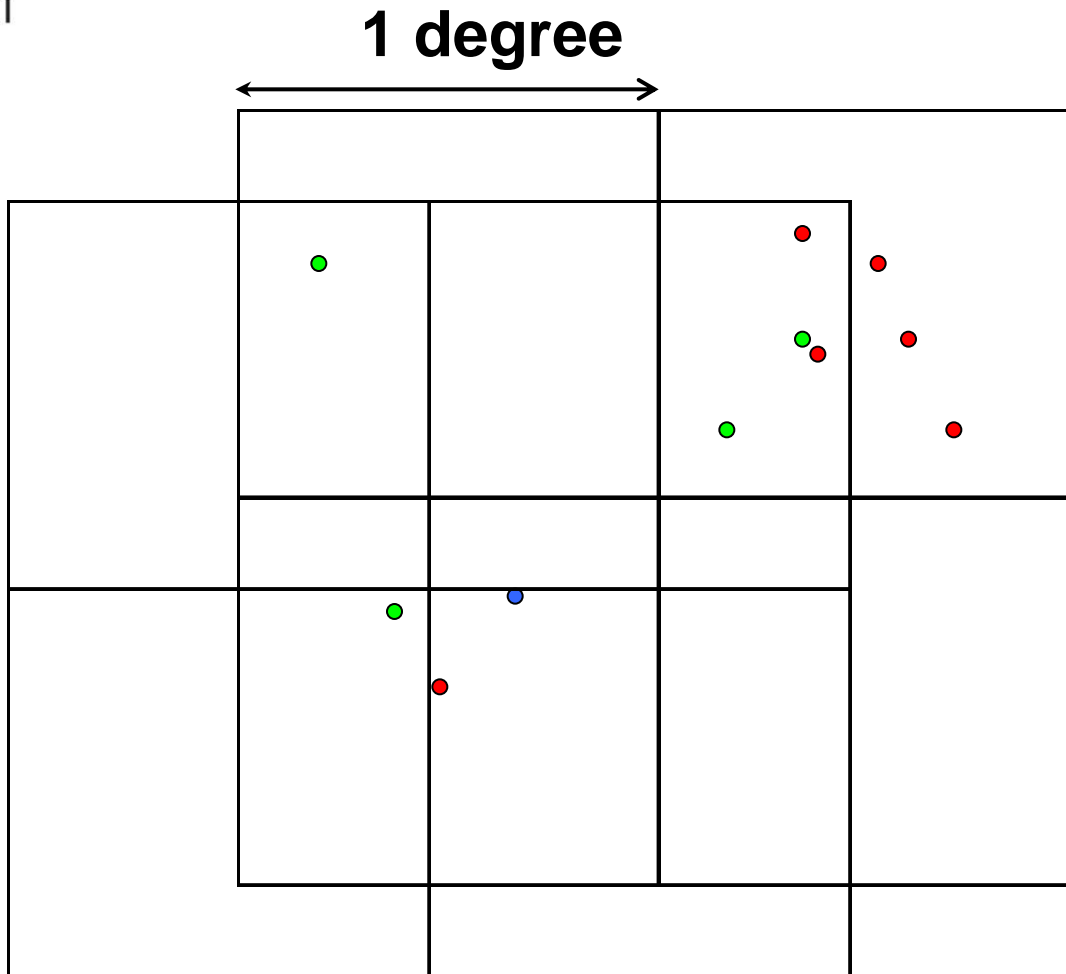


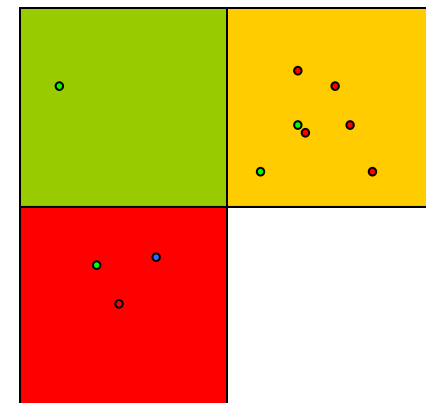
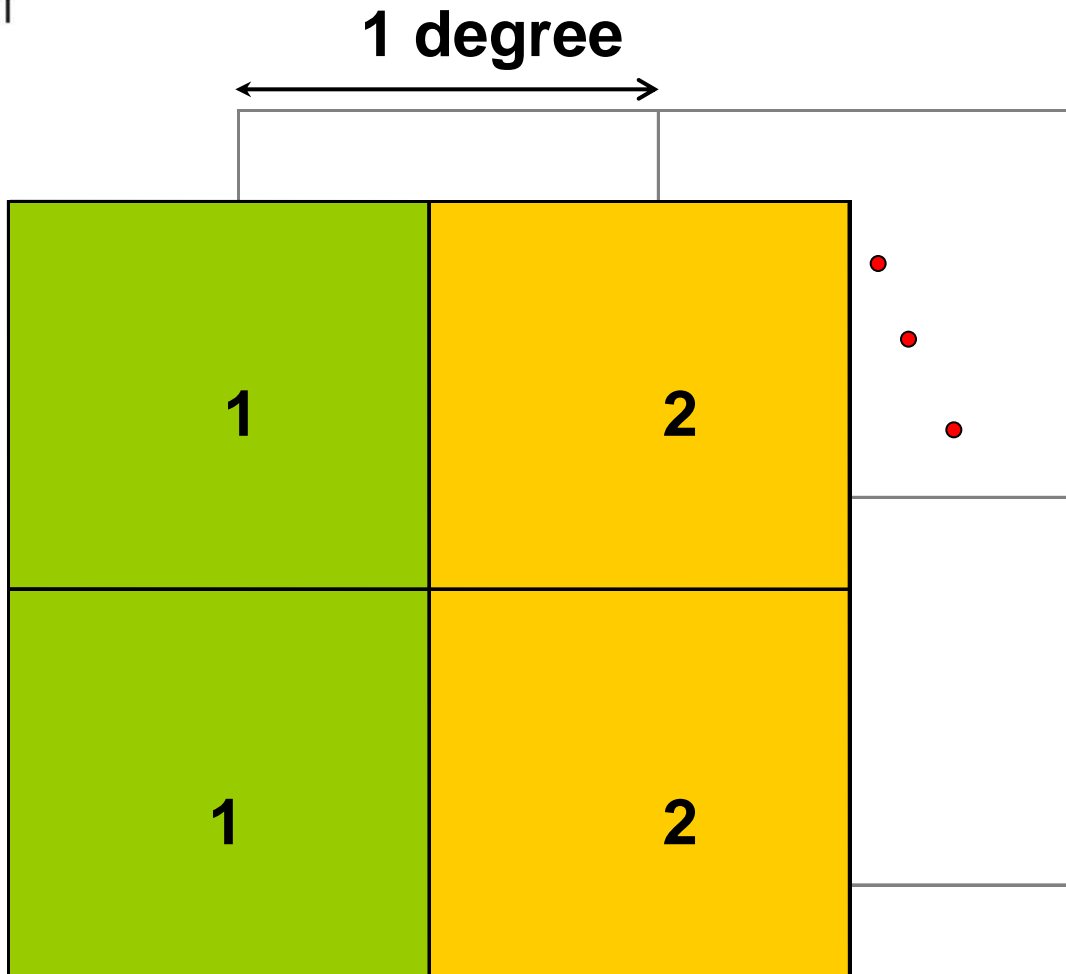


Diversity question...

Where can the highest *Vasconcellea* diversity be found?







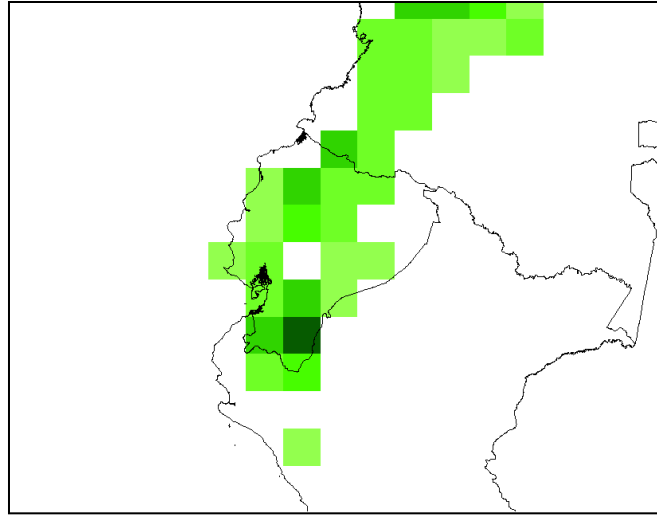
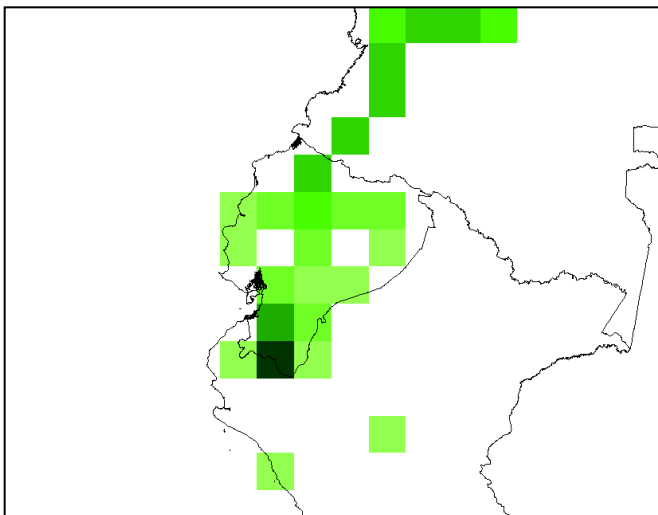
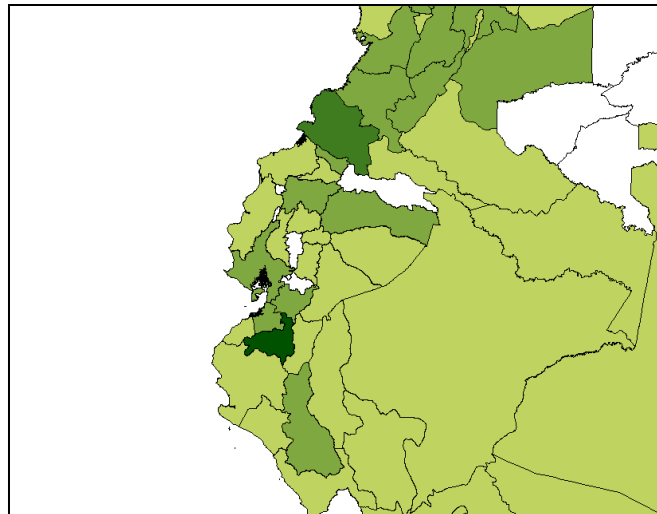
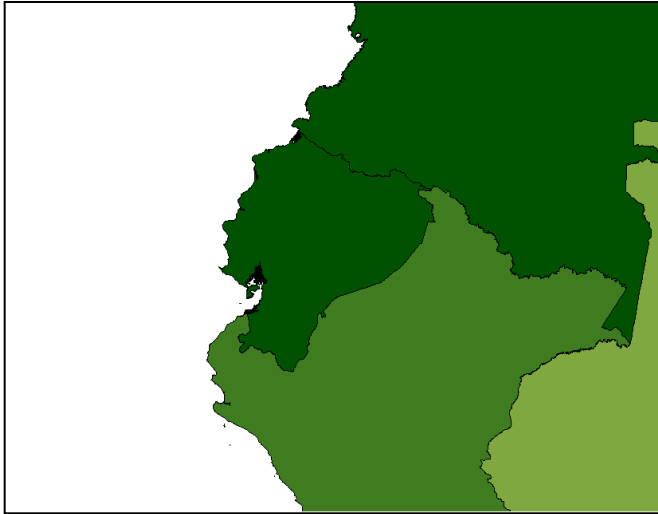
Diversity question...

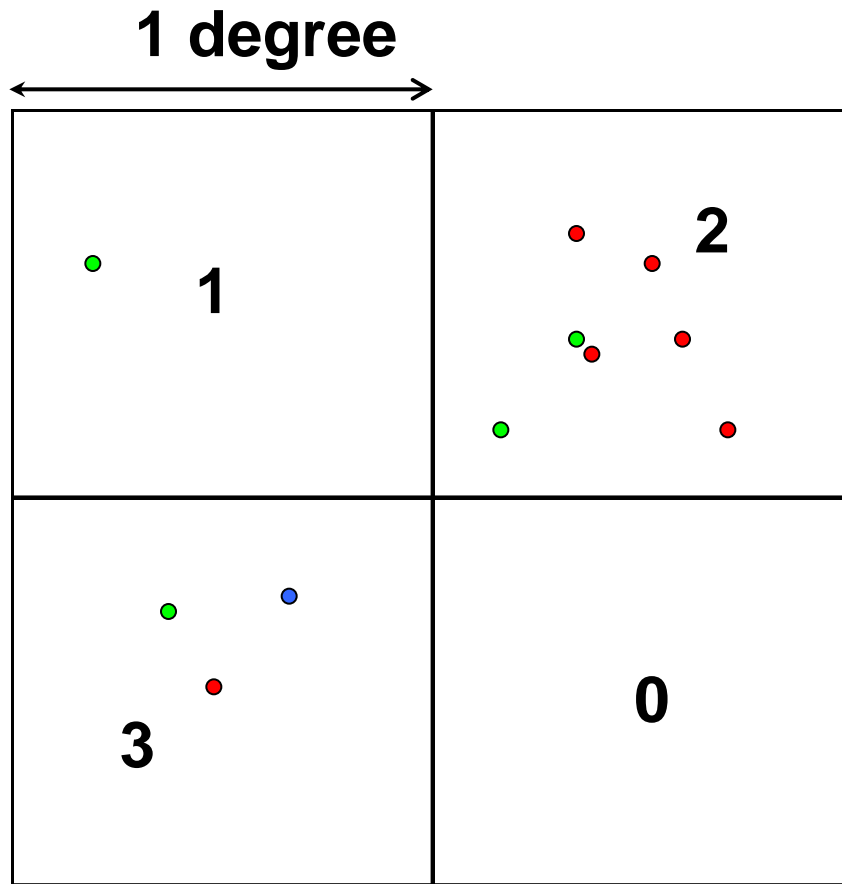
Where can the highest *Vasconcellea* diversity be found?

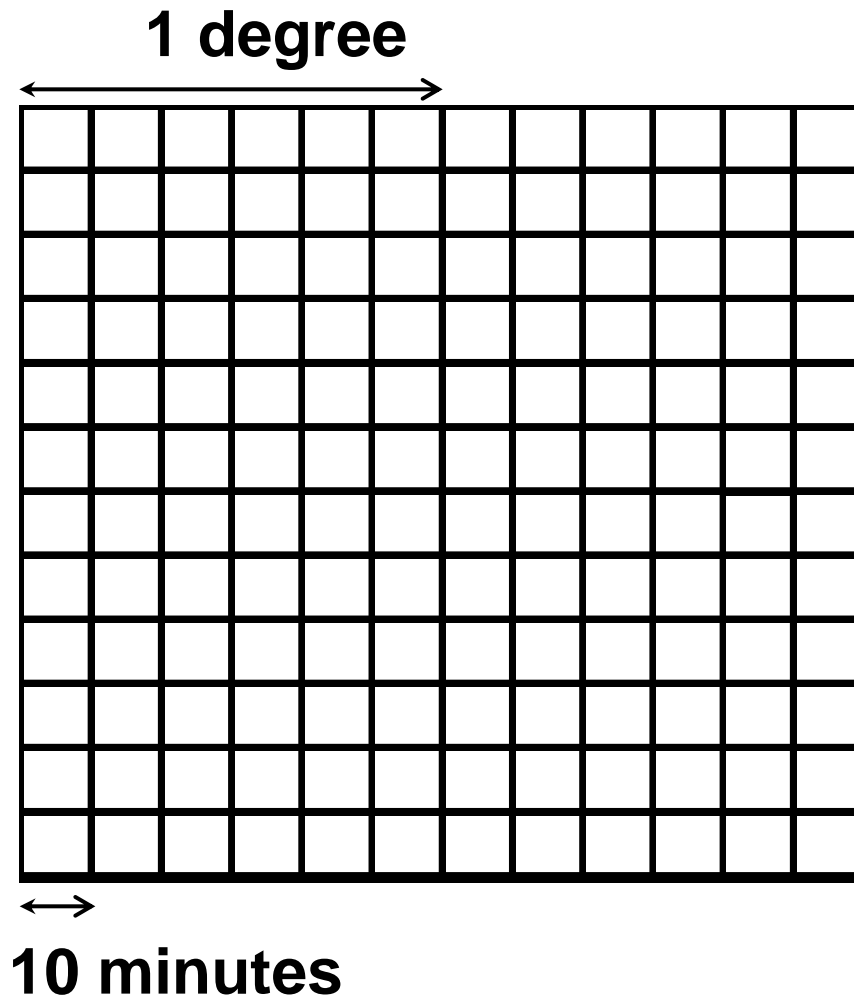


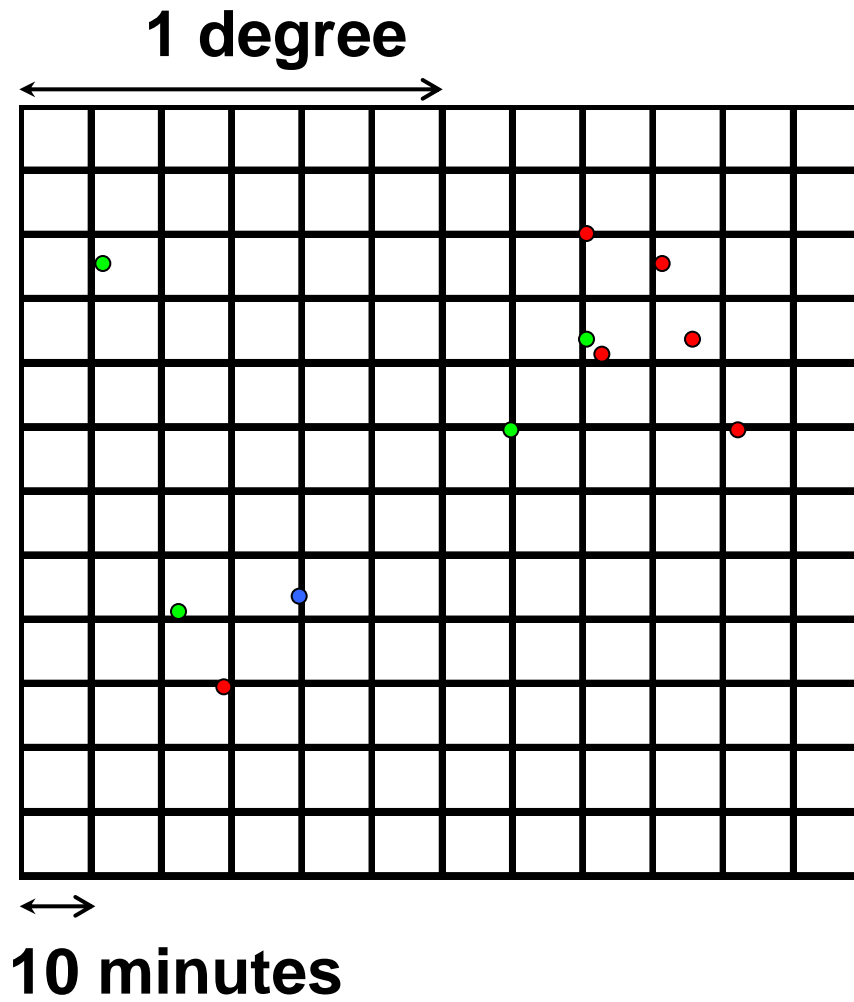
Diversity question...

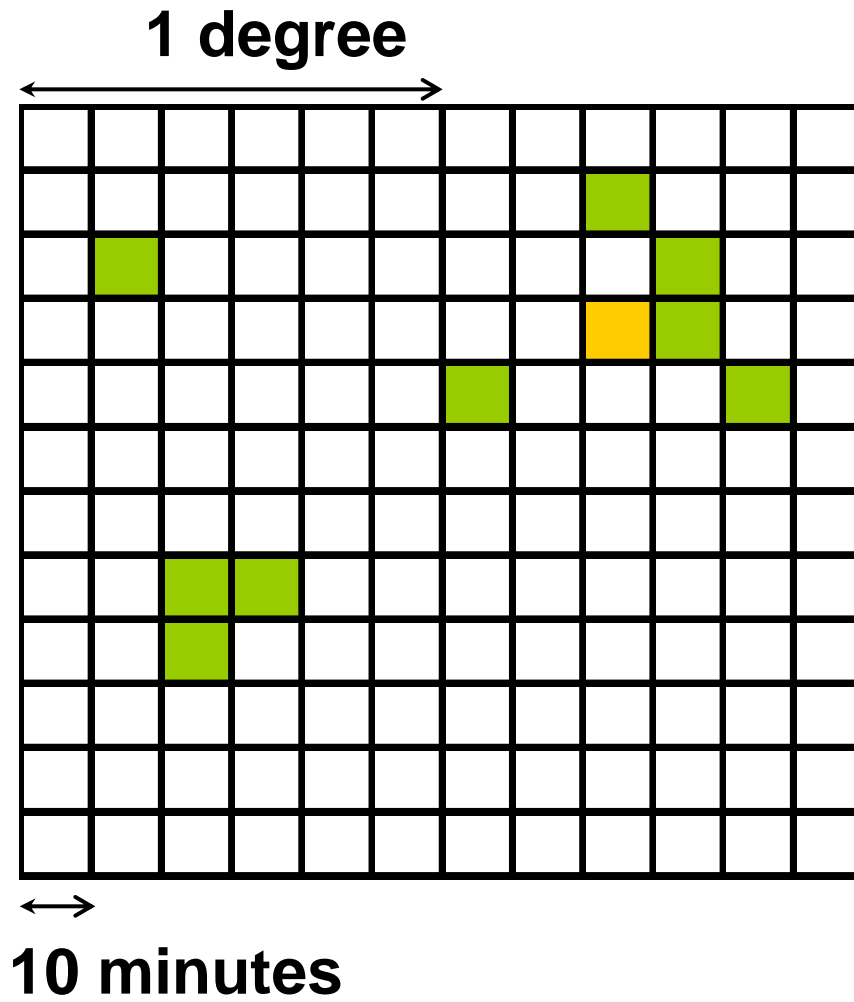
Where can the highest *Vasconcellea* diversity be found?





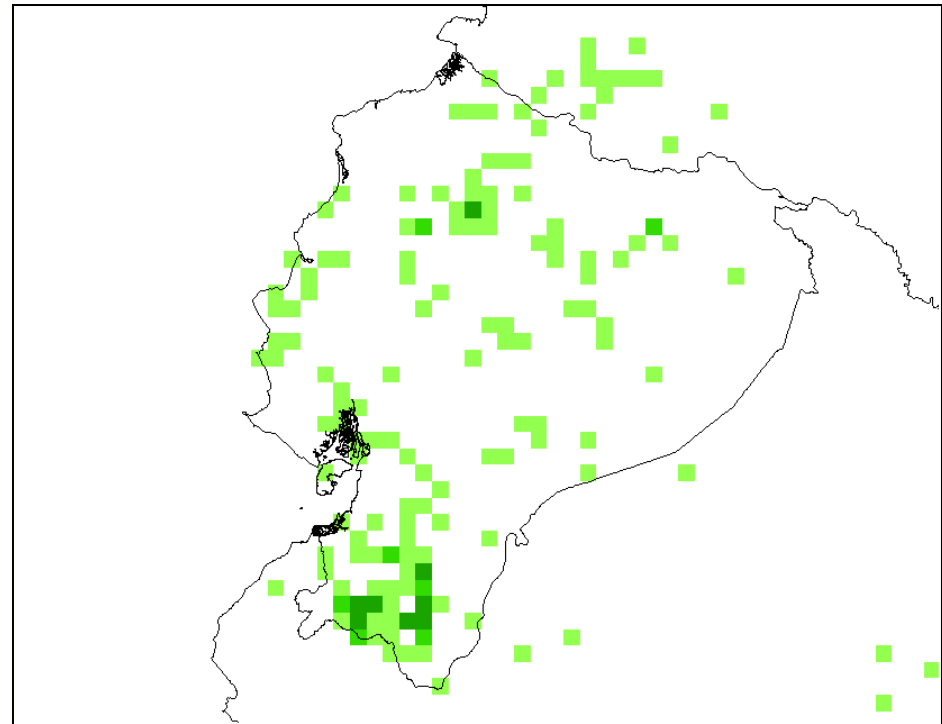
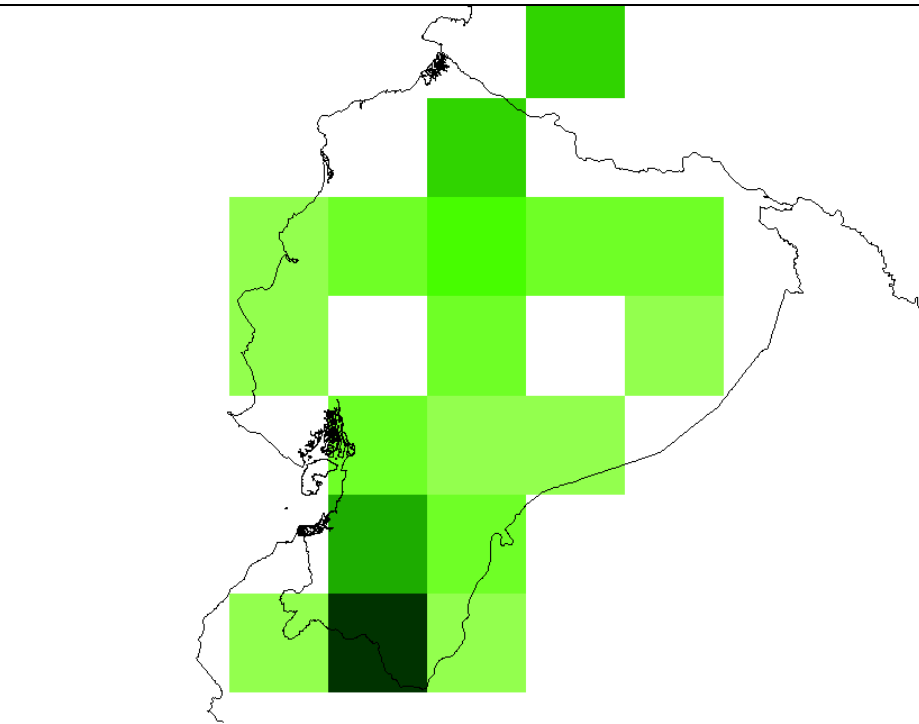


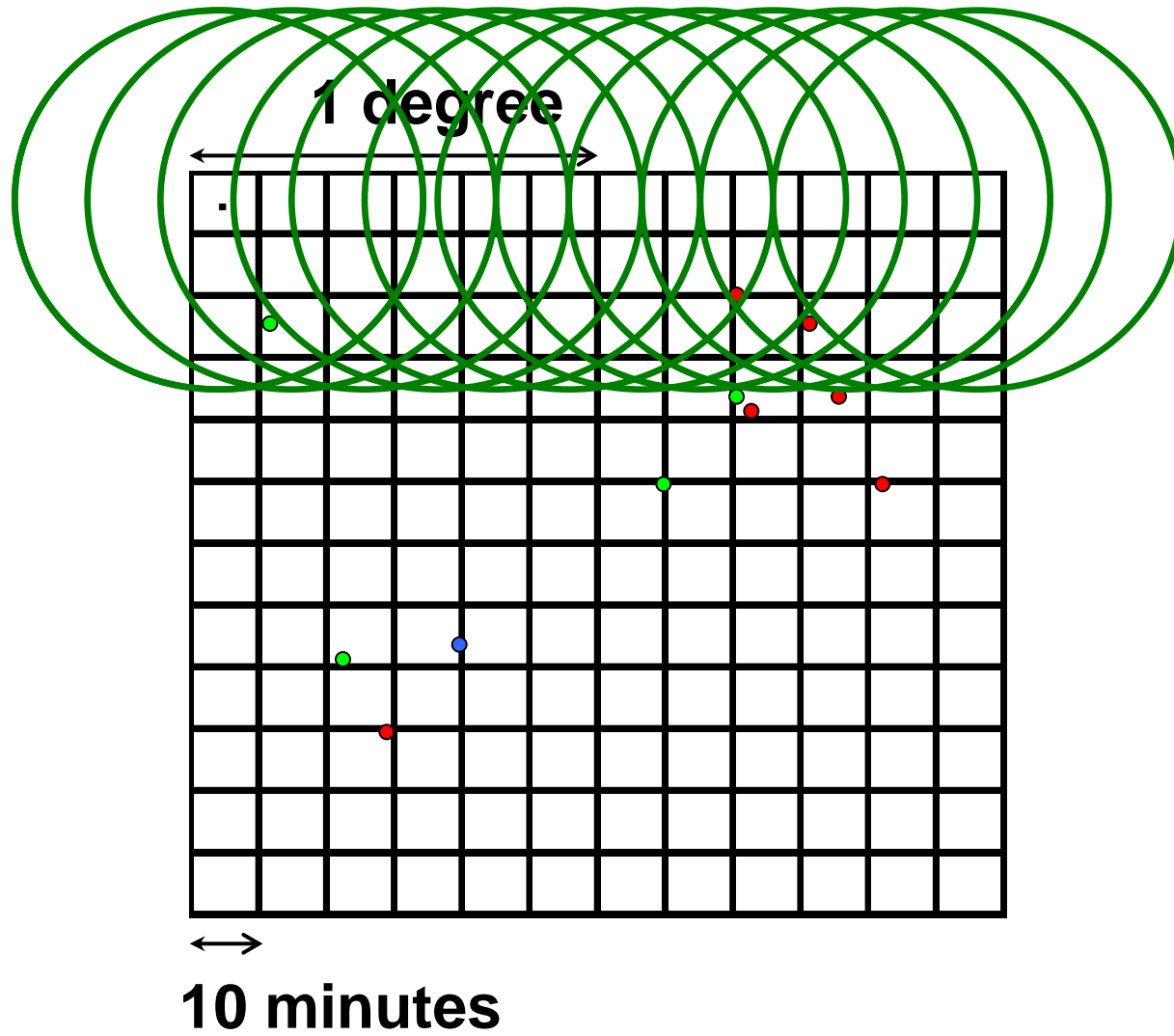


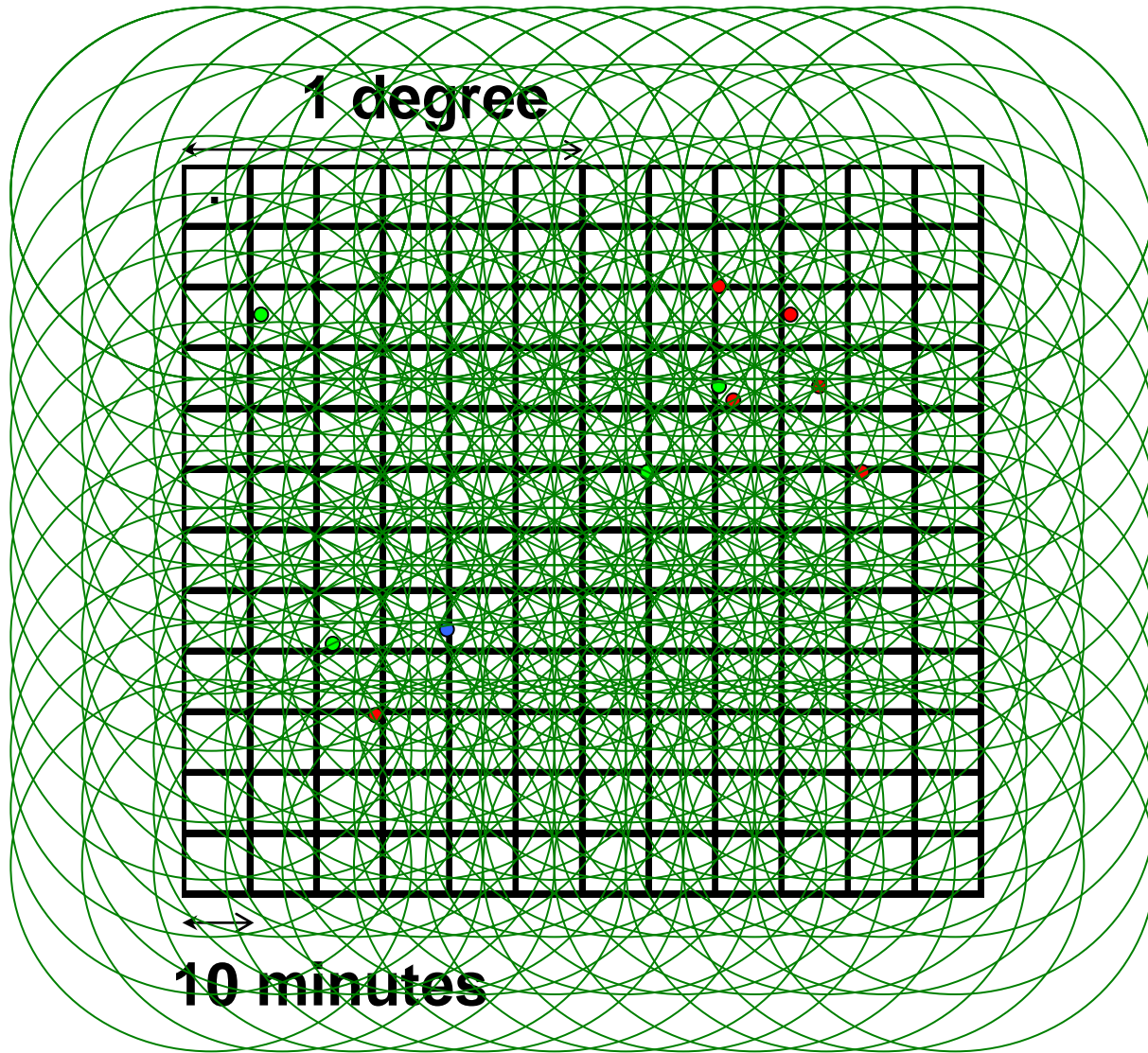


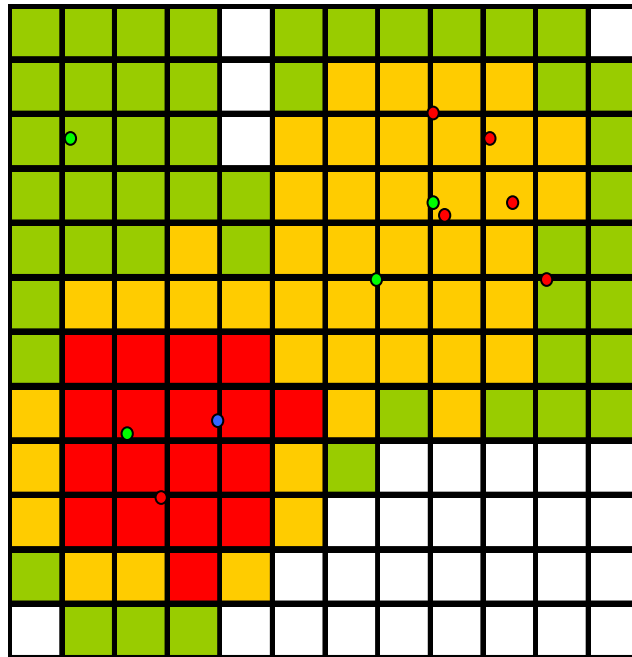
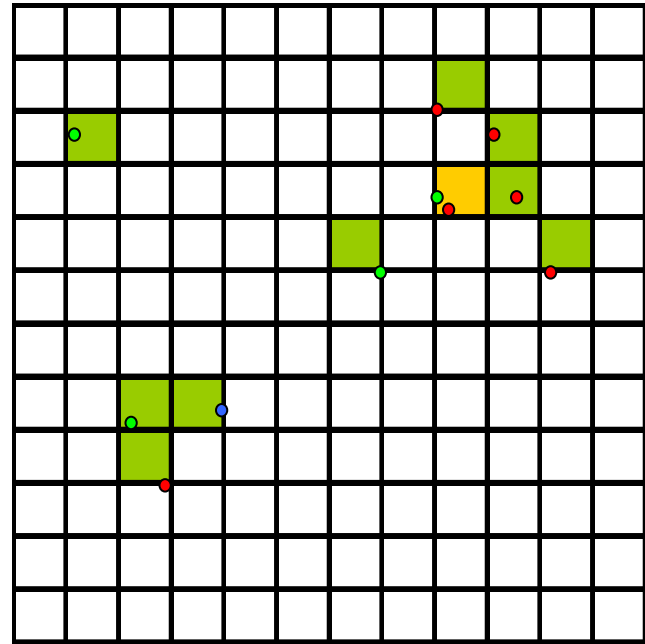
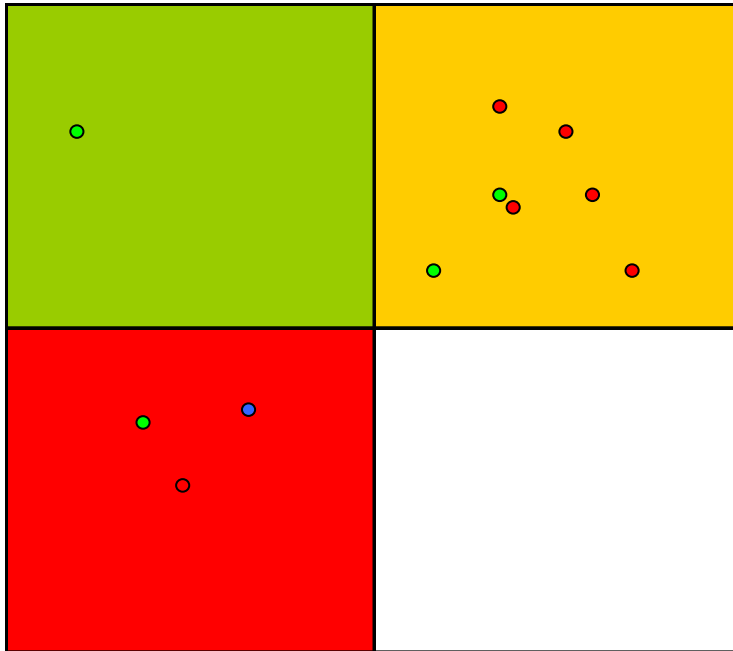
Diversity question...

Where can the highest *Vasconcellea* diversity be found?



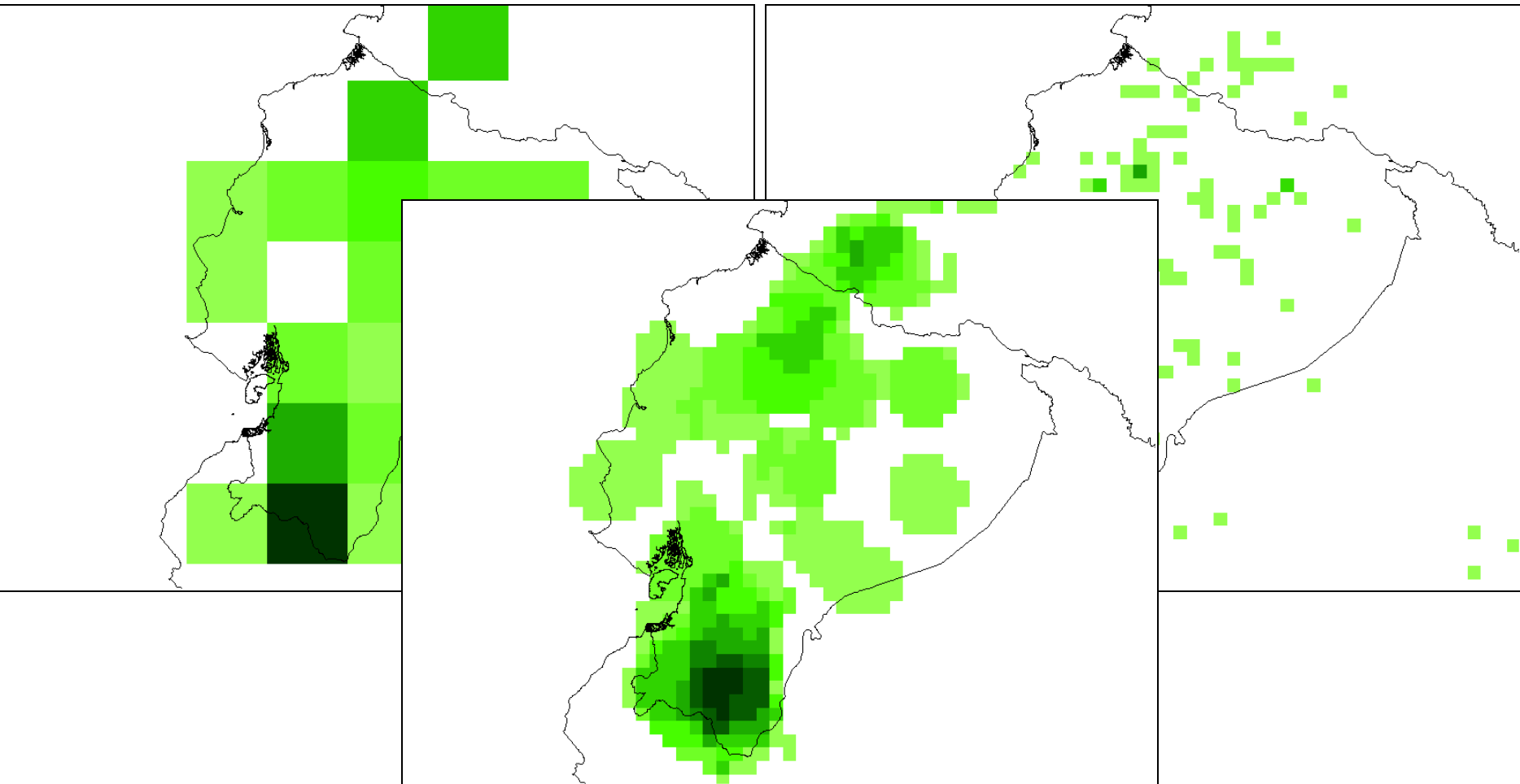






Diversity question...

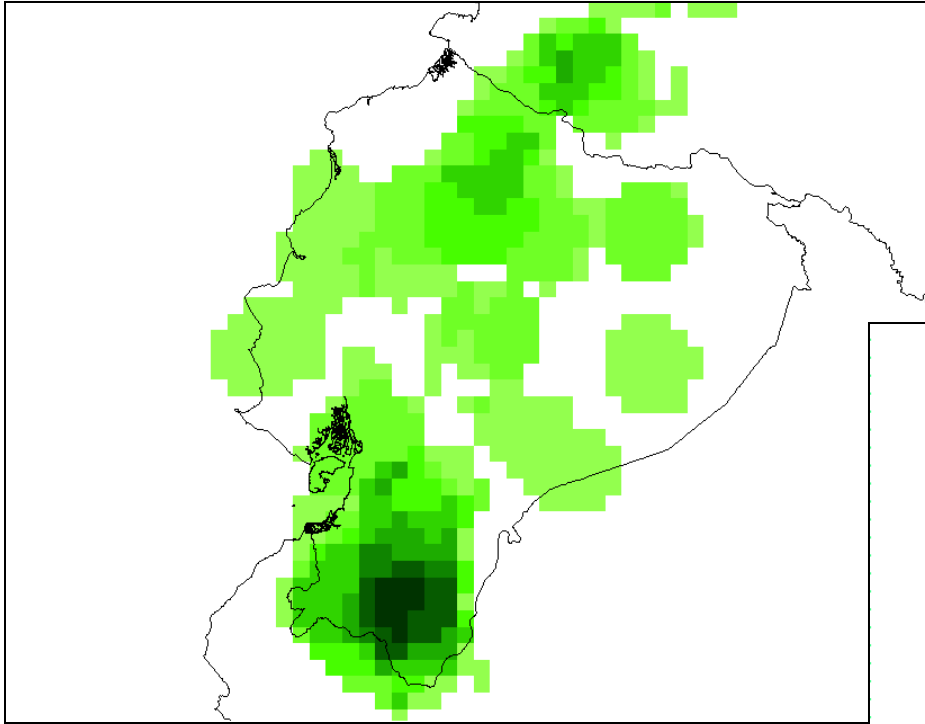
Where can the highest *Vasconcellea* diversity be found?



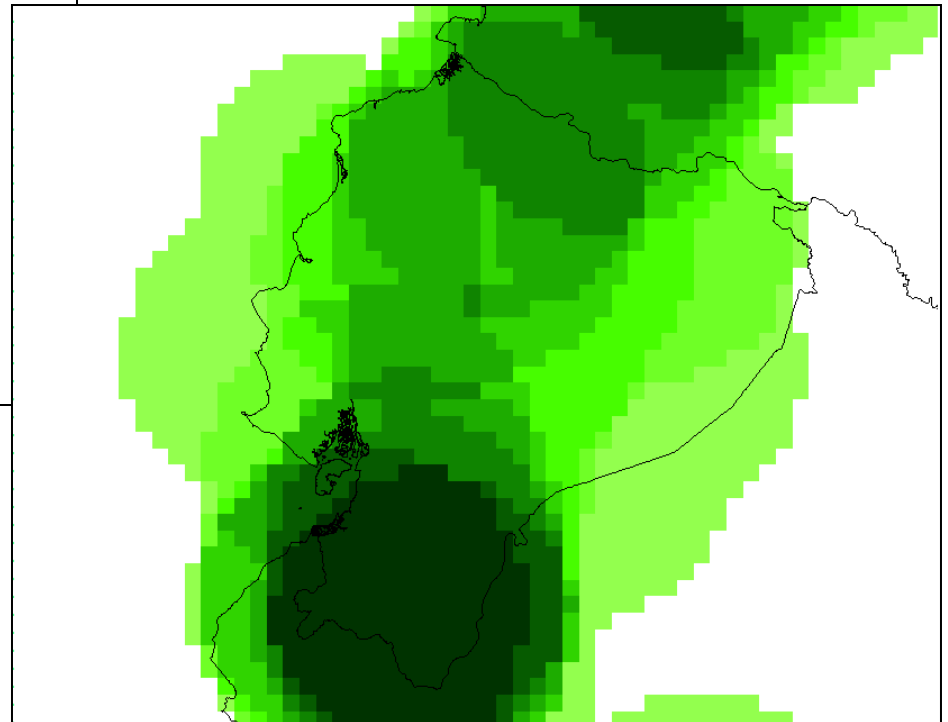
Diversity question...

Where can the highest *Vasconcellea* diversity be found?

Optimal circle radius?



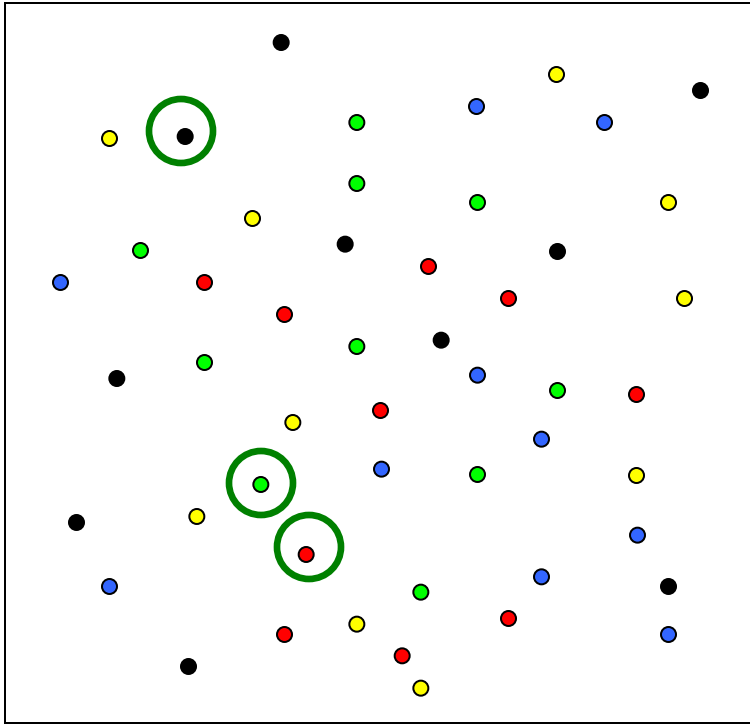
Cell: 10 minutes
Circle: 1 degree



Cell: 10 minutes
Circle: 3 degrees

Sampling bias

50 observations

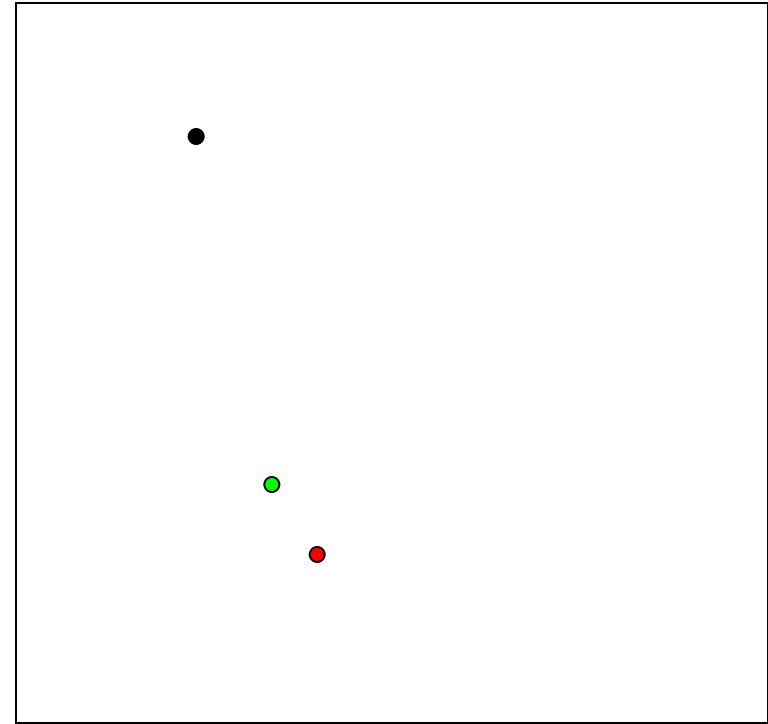


5

Richness

Possible solutions

3 observations



3

Reduce resolution (bigger cells)

Prediction of missing distribution (modelling)

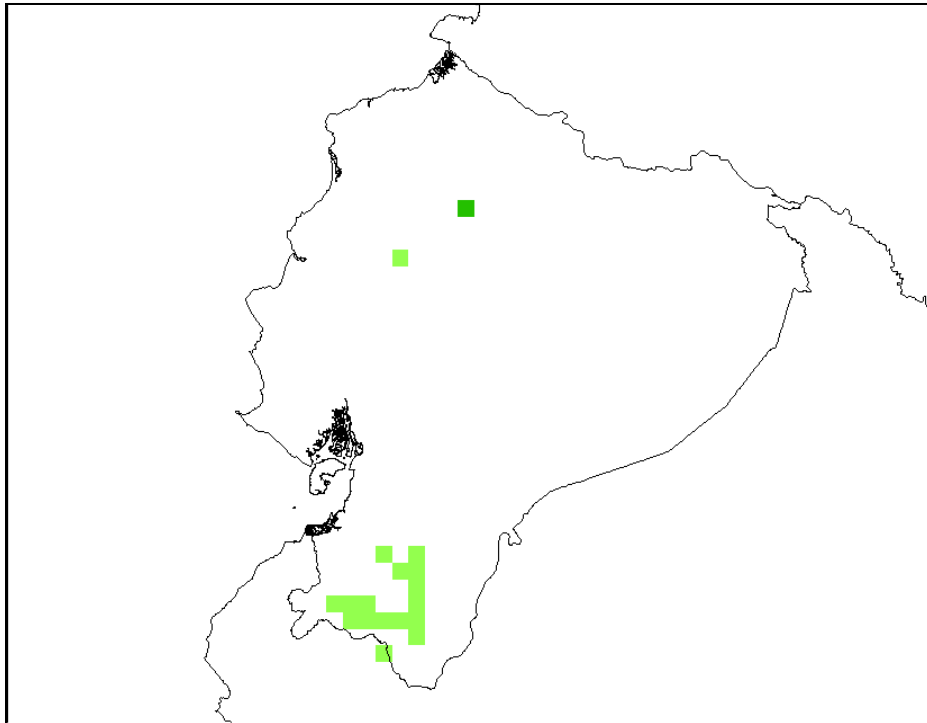
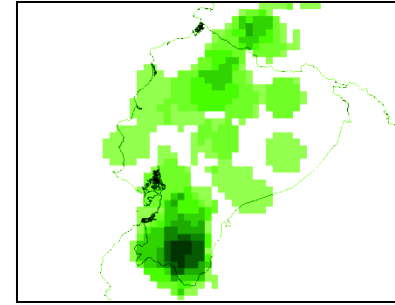
Rarefaction

The rarefaction technique re-calculates the number of classes (species in this case) that would have been observed given a number of observations that is specified by the user (based on hypothetical subsampling)

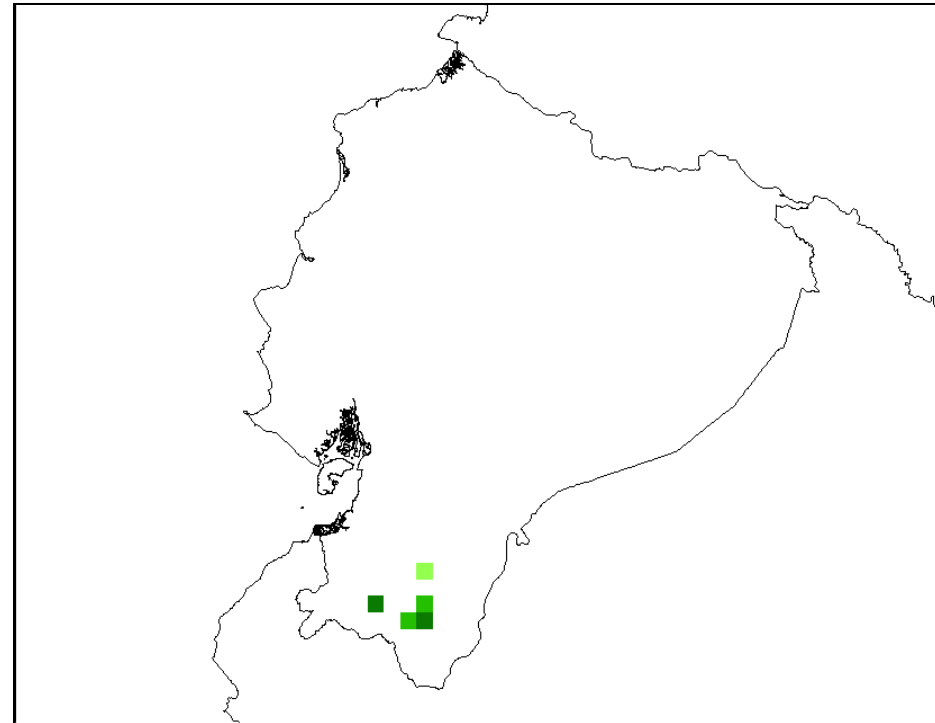
User defined number of observations?	Final cell value	Number of cells	Final results
Too low (e.g. 5)	Max richness defined by defined number of observations (e.g. 5)	Many	Loss of information – loss of detail of final cell value
Too high (e.g. 15)	High	Only those cells with a number of observations above the selected number will be included (e.g. 15)	Loss of information – loss of cell value

Sampling bias

Rarefaction



reduction to 3 observations



reduction to 10 observations

Applicability in case of limited number of observations?

Estimators of Richness

Chao 1 $S_1 = S_{OBS} + (a^2 / 2b)$

Chao 2 $S_{Chao2} = S_{obs} + \frac{Q_1^2}{2Q_2}$

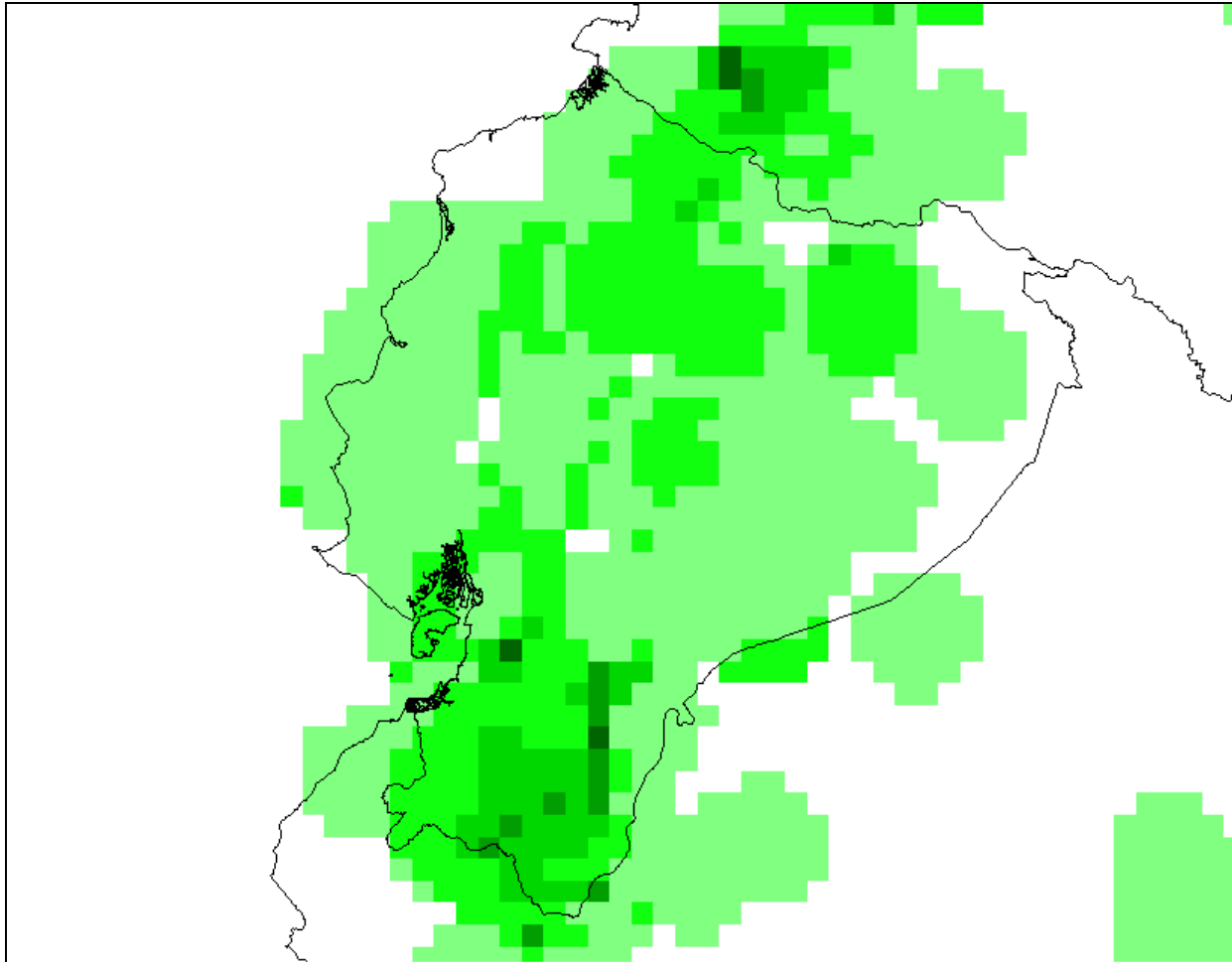
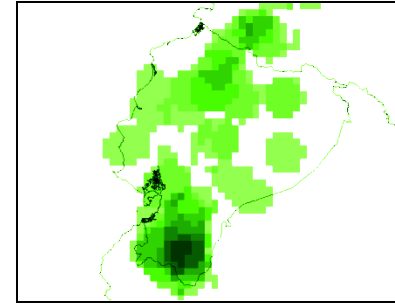
Chao 1 Corrected $S_{Chao1} = S_{obs} + \frac{F_1^2}{2(F_2 + 1)} - \frac{F_1 F_2}{2(F_2 + 1)^2}$

... (cf DIVA manual)

Applicability in case of limited number of observations?

Sampling bias

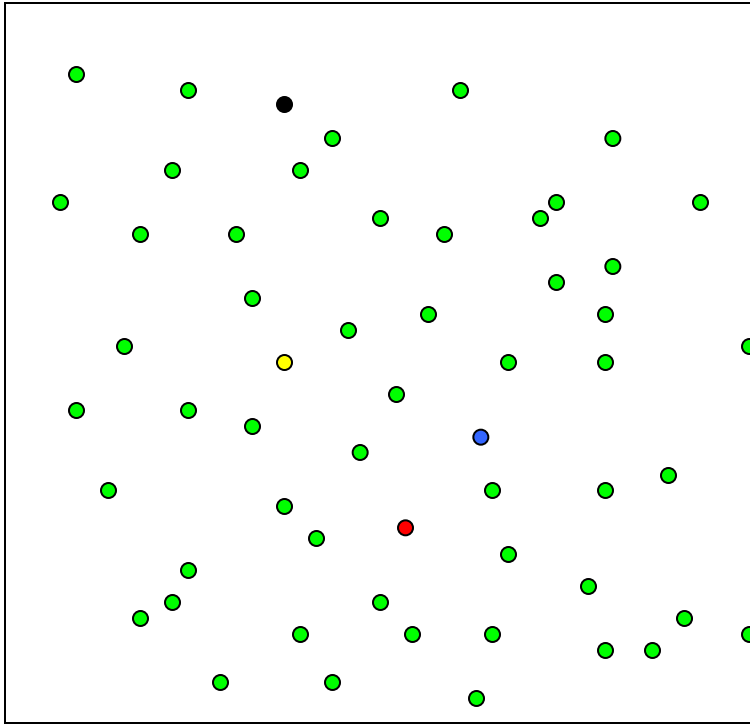
Estimators of Richness



Chao 1 Corrected

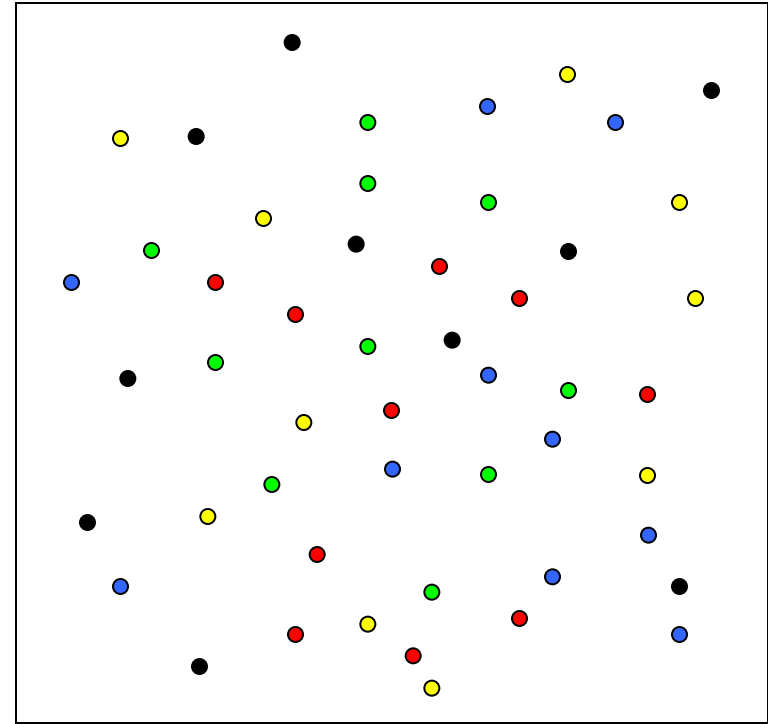
Other measurements of diversity

50 observations



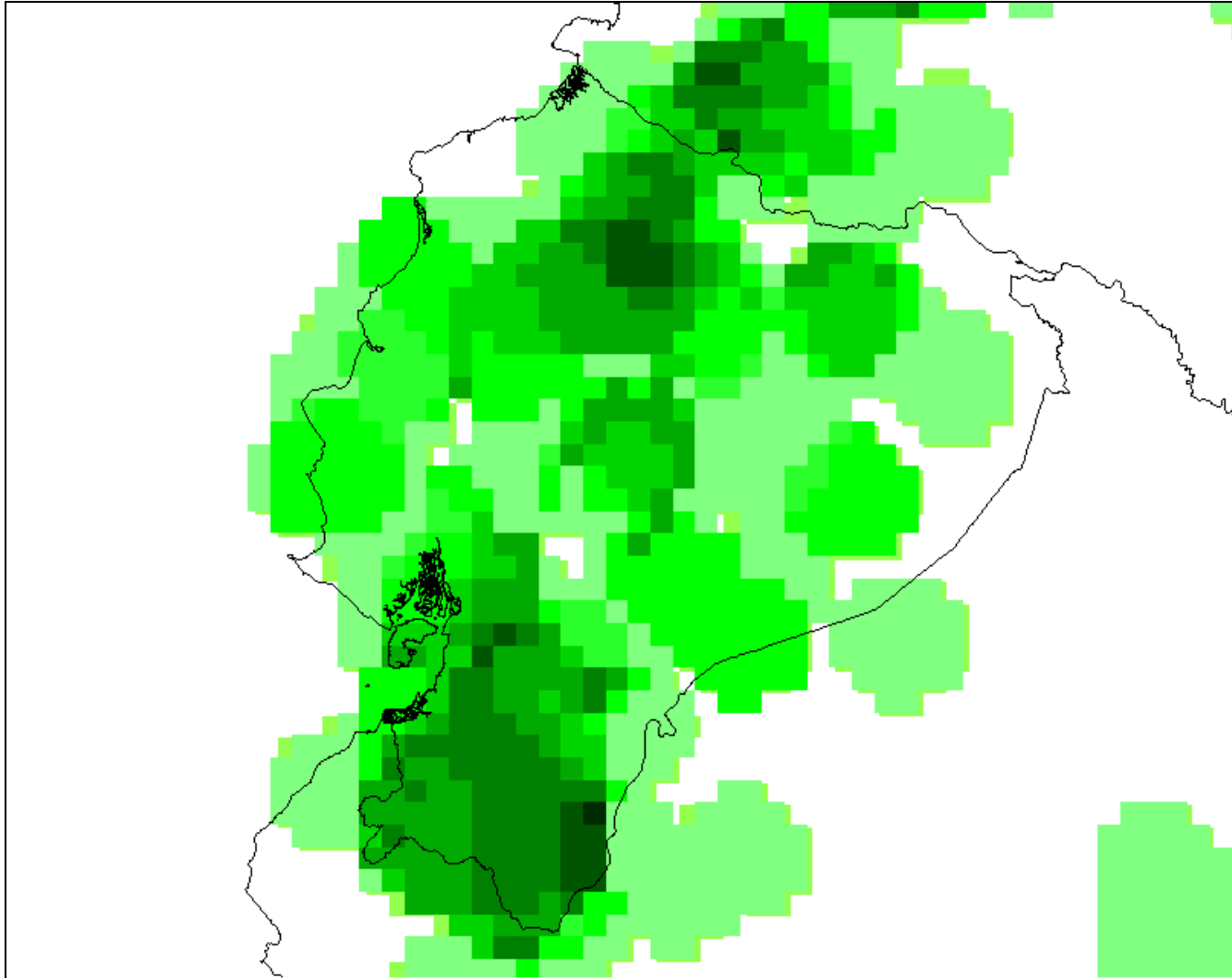
5

Richness



5

Other measurements of diversity



Shannon diversity

Quantitative data

- Range
- Range / Mean
- Standard deviation
- Coefficient of variance

Molecular data

- Observed allele number
- Heterozygosity (external software)
- Effective allele number (external software)

Going beyond the cell level

Alpha versus beta diversity

- **Alpha diversity (α -diversity)** is the biodiversity within a particular area, community or ecosystem and is usually expressed by the number of species (*i.e.*, species richness) in that ecosystem
- **Beta diversity (β -diversity)** is a measure of biodiversity which works by comparing the species diversity between ecosystems or along environmental gradients.
- **Gamma diversity (γ -diversity)** is a measure of the overall diversity for the different ecosystems within a region

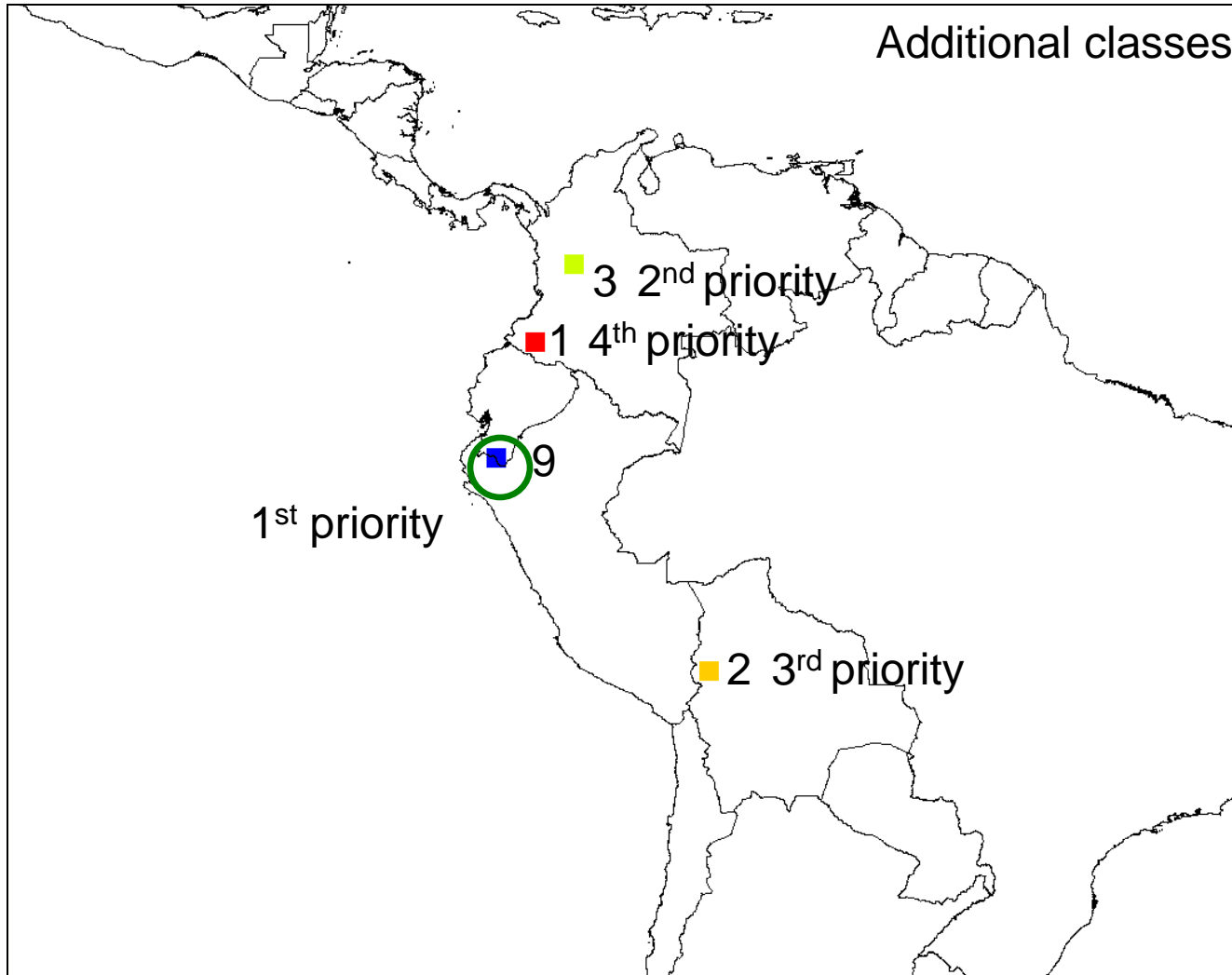
Whittaker's measure

$$\beta = (S1 - c) + (S2 - c)$$

$$\beta = \frac{S}{\bar{\alpha}} \text{ or } \beta = \frac{S}{\bar{\alpha}} - 1$$

$$\gamma = S1 + S2 - c$$

Reserve Selection



Priority areas for in situ conservation?



Thanks