Disturbance – Human Footprint gradient (%) All Alberta

Data summary

- Vegetation data PA from GitHub
  - 2504 unique ID (ID = protocol + site + year)

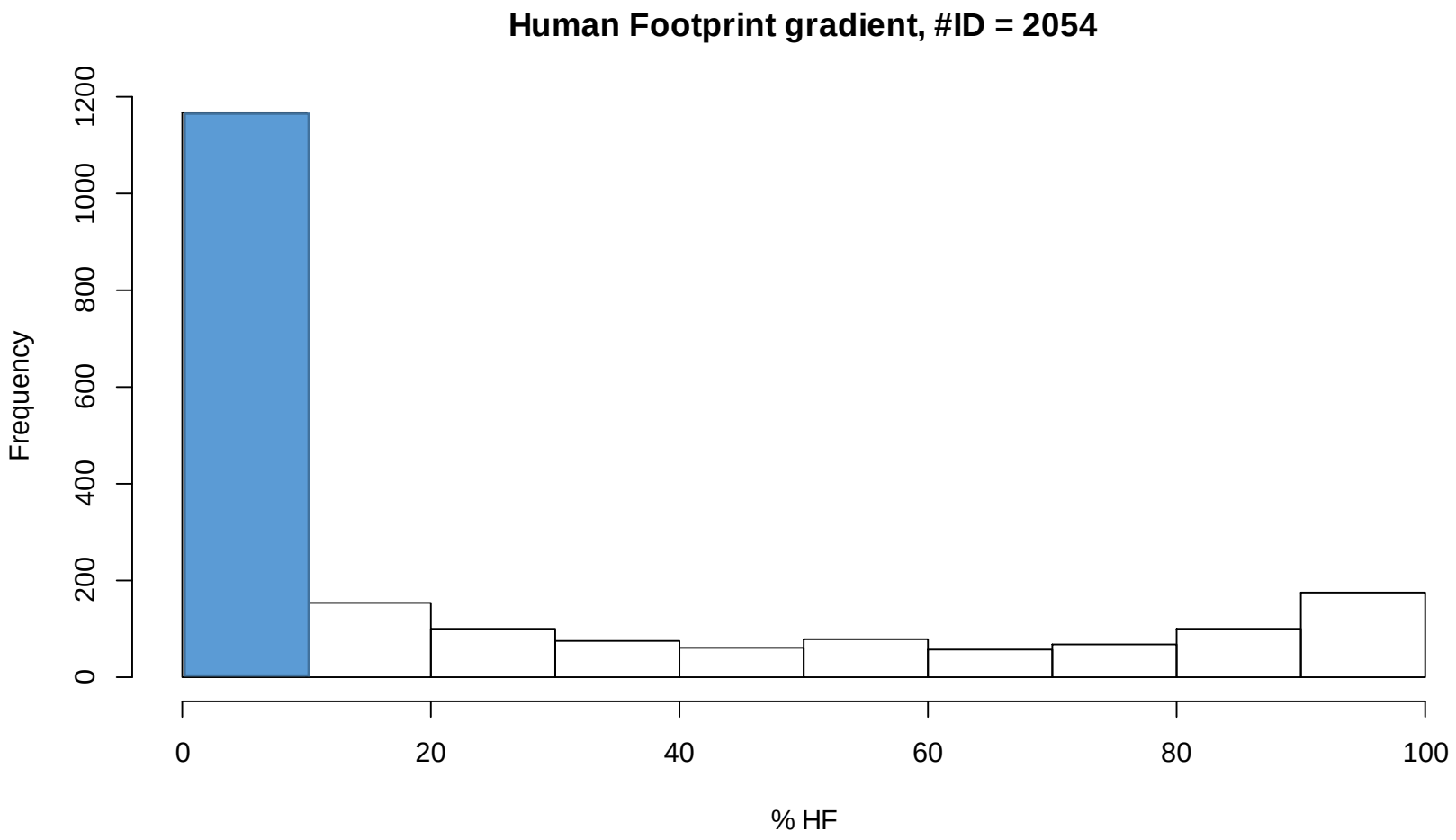
- Human Footprint (HF) % data from GitHub
  - 1866 unique ID

- 277 ID of vegetation data are not in HF data :
  - Not equal to 2504-1866 = 188 ID. Why ? Because HF data with value = 0% for terrestrial ID are not present (obvious for example because all Canadian Shield terrestrial ID are not in the HF dataset).
  - I assumed that all ID of vegetation data without HF data have HF data = 0%

- Combining this way vegetation and HF data, I obtained : 2054 ID for the final dataset

Data summary

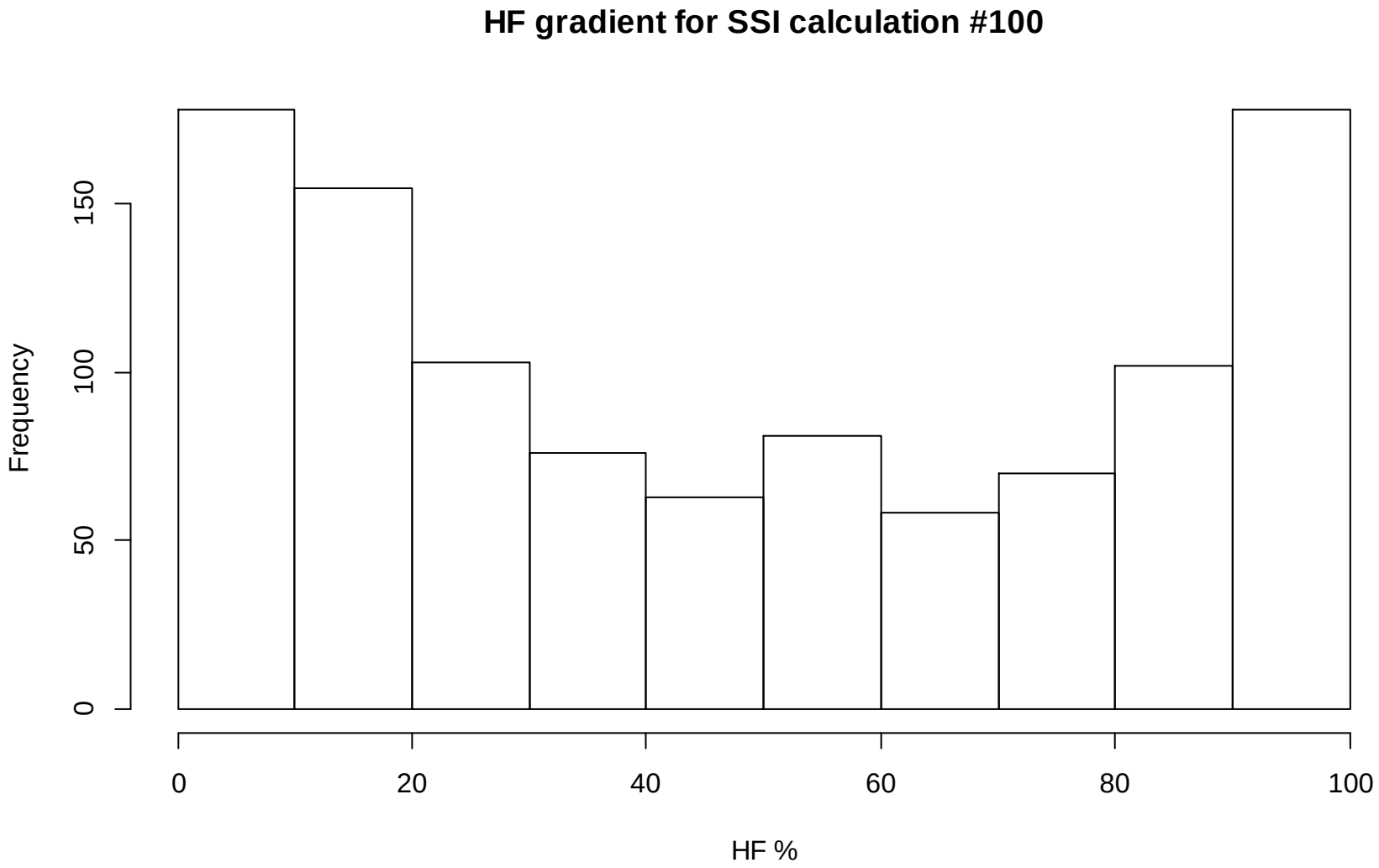
- Distribution of the number of samplings in 10 classes of HF are very uneven:



Applied randomization process previously showed (see previous ppt). #run = 100

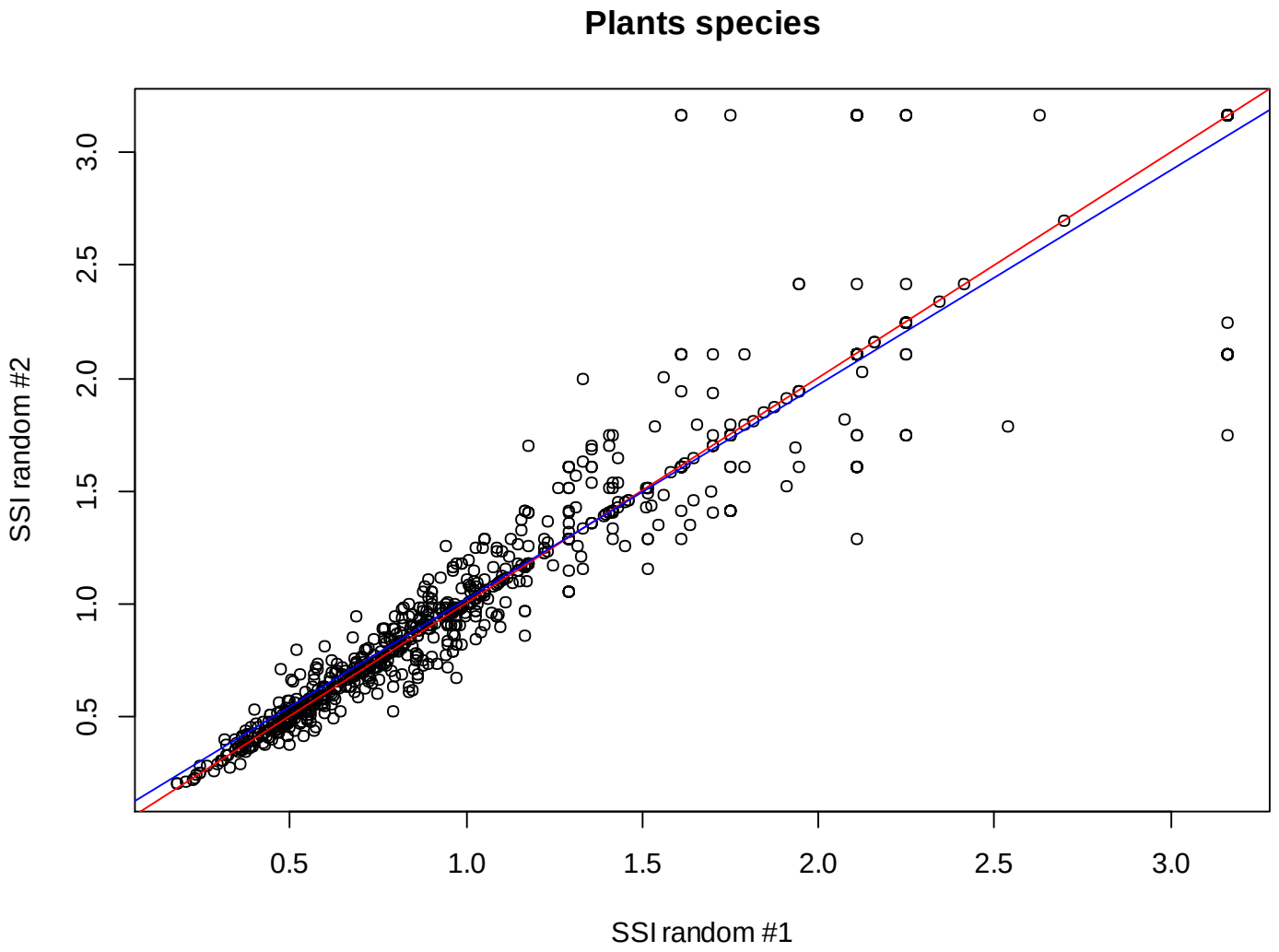
Obtained 10 categories of HF % where I tried to have each time the same number of samplings per category (see previous ppt)

- Distribution of HF % used for one (over 100) calculation of SSI values:



Calculate SSI on the 10 categories for each random dataset - Plants

- Look at correlations between the calculated SSI (n = 100)
  - mean correlation values = 0.94 ± 0.01



Good overall correlations

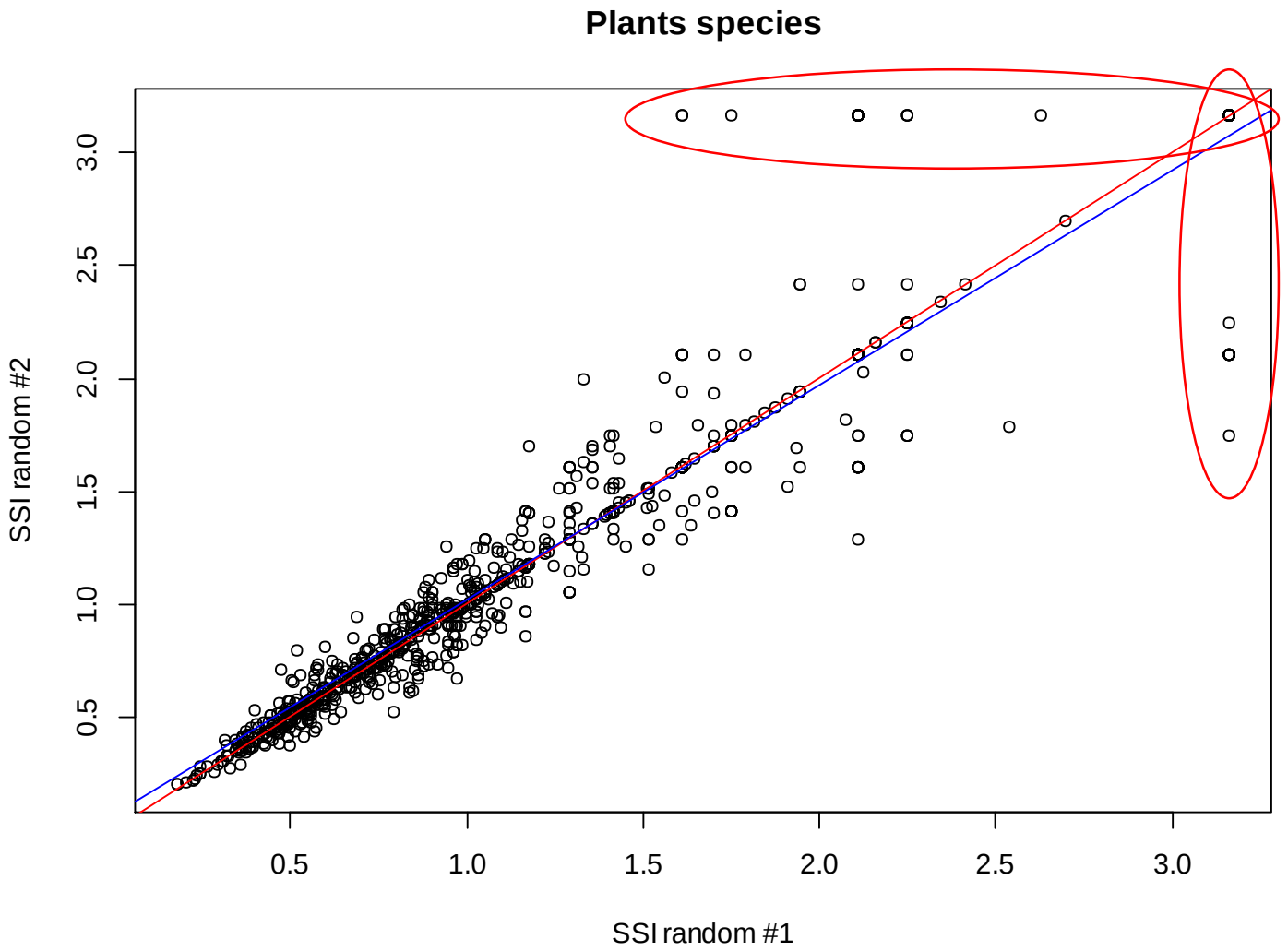
Red line = perfect correlation

Blue line = linear model of actual relationship between two random set of SSI calculation

→ Take one of the random calculation of SSI for next analyses

Calculate SSI on the 10 categories for each random dataset - Plants

- Look at correlations between the calculated SSI (n = 100)
  - mean correlation values = 0.94 ± 0.01

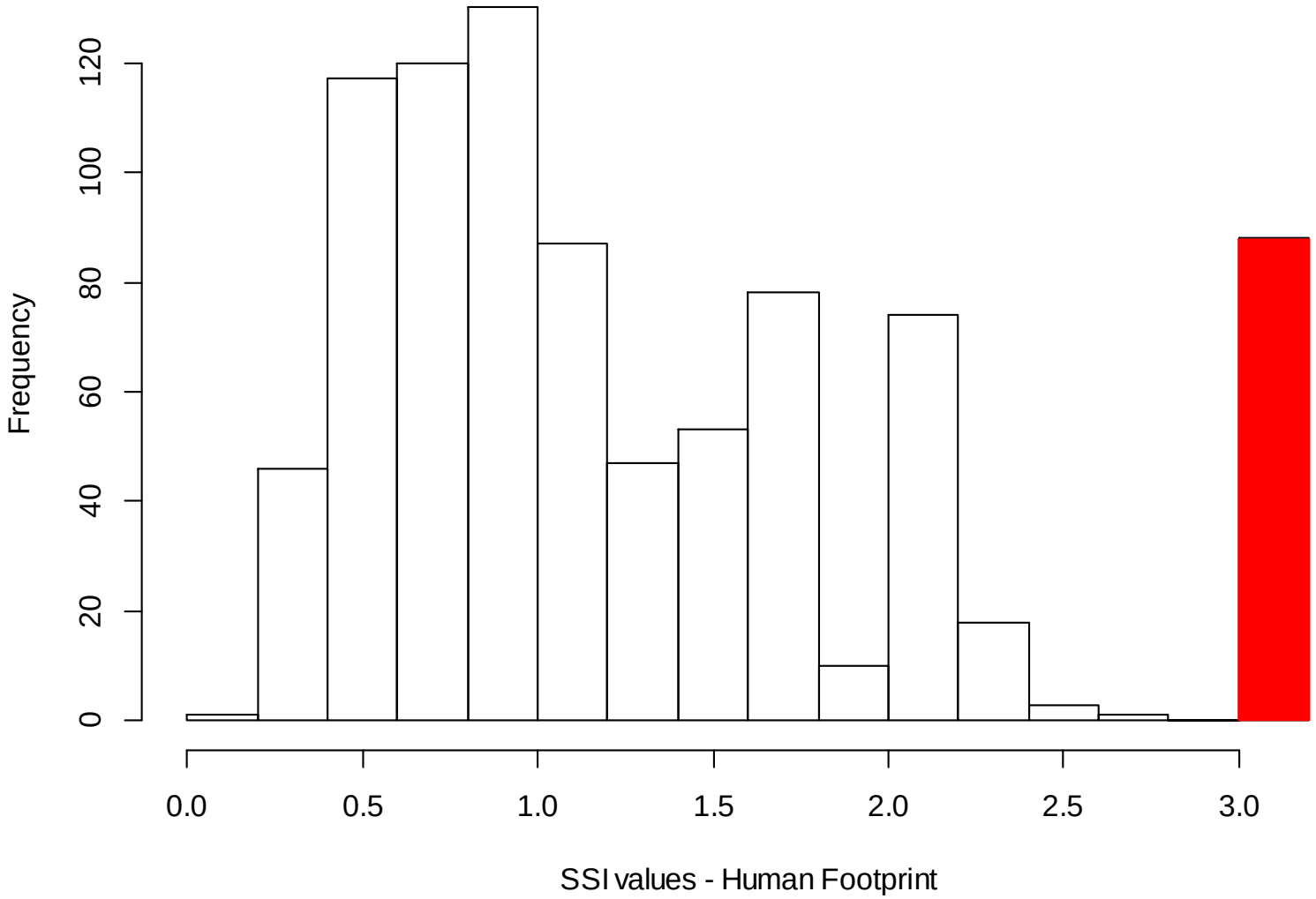


Red ellipses show outliers values that should be deleted :

Variance of SSI values for these species are too high

Expected to be rare species with not reliable SSI calculation.

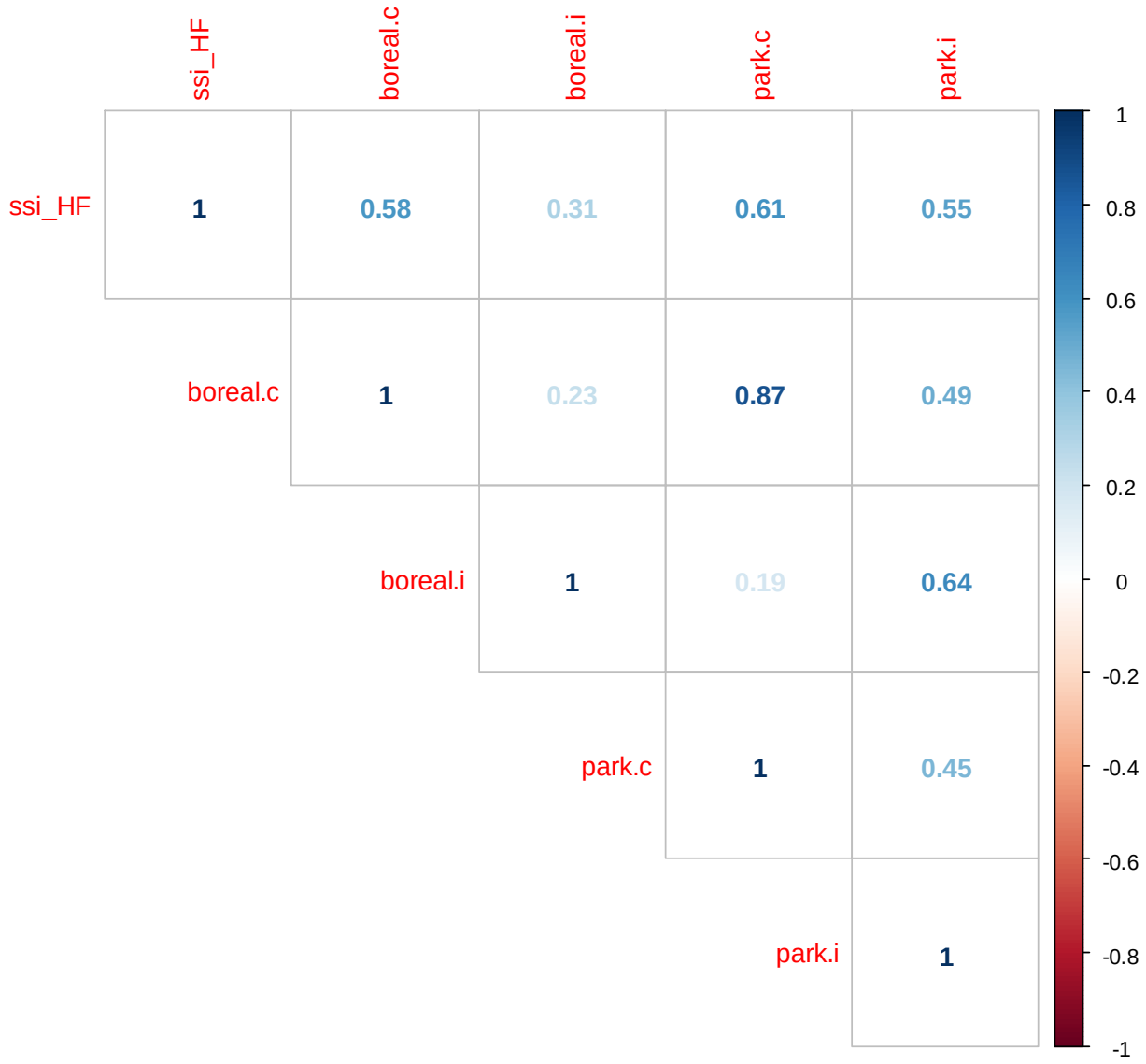
Distribution of SSI values (randomly chosen from the hundred calculated) - Plants



Red Bar = outliers SSI values (the one in the red ellipses in the previous slide) that are deleted for next analyses (i.e. removing these species of the communities)

88 species, mostly present on 1 or 2 ID (max 4)

Comparison of SSI values with c-scores and intactness scores - Plants

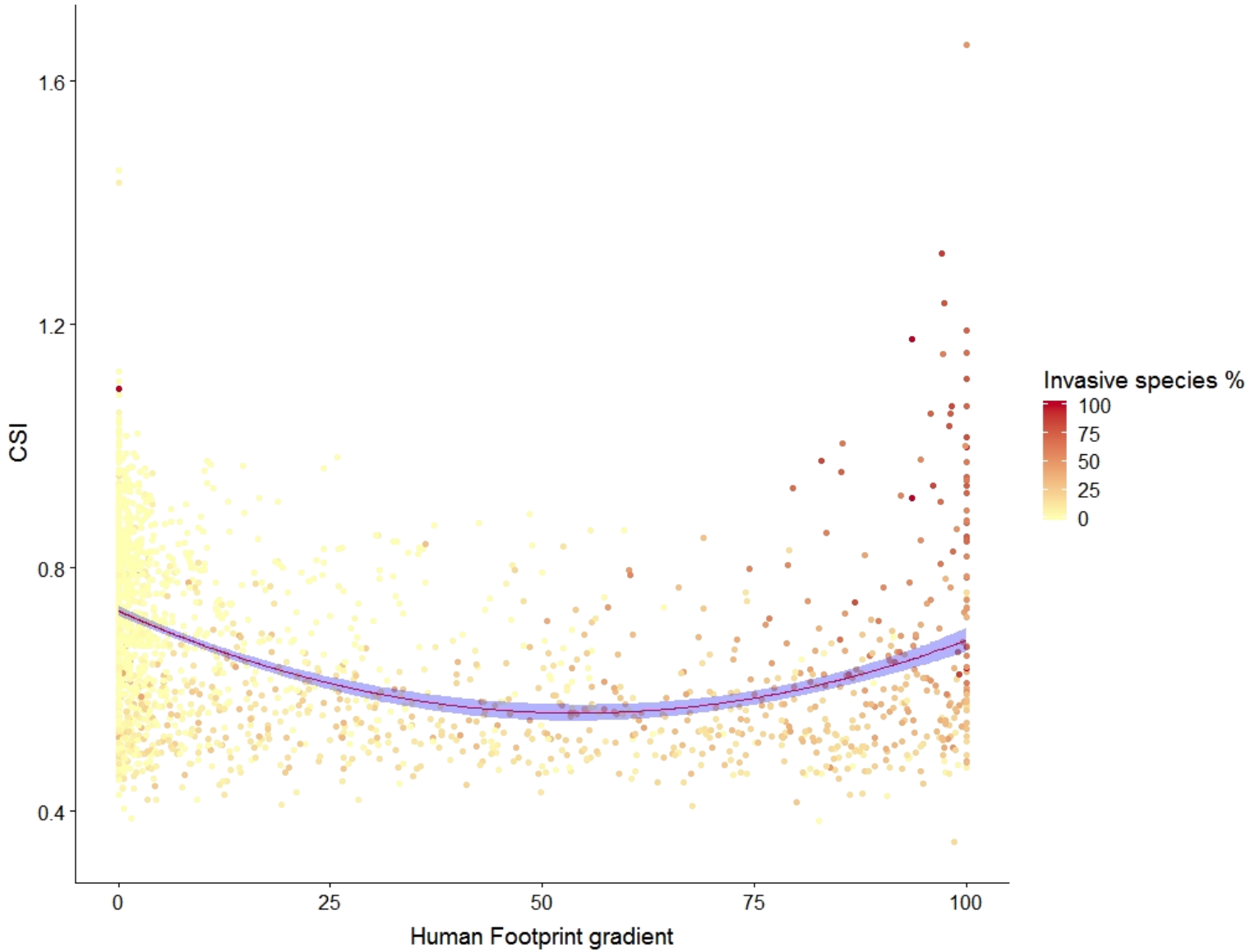


SSI good indicator of conservatism ?

Easy to implement
Not based on expert opinion

Calculation of CSI values and plot along the HF gradient + Invasive species proportion

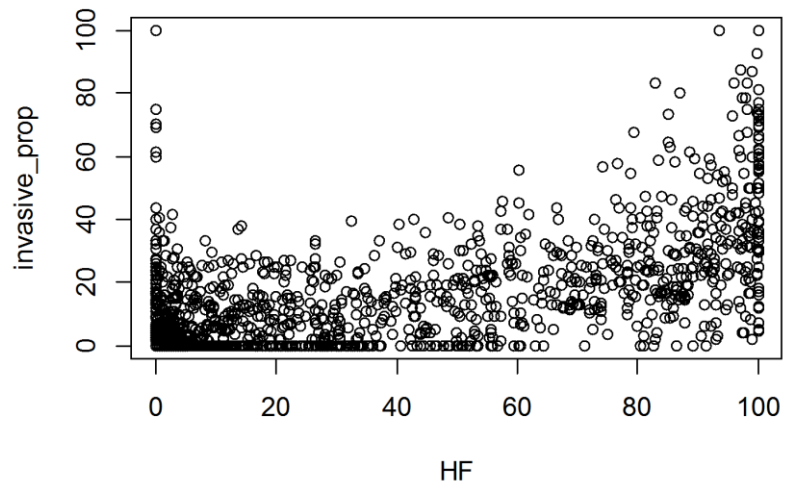
- CSI = Community specialization index = mean of SSI present in each community



Interpretation:

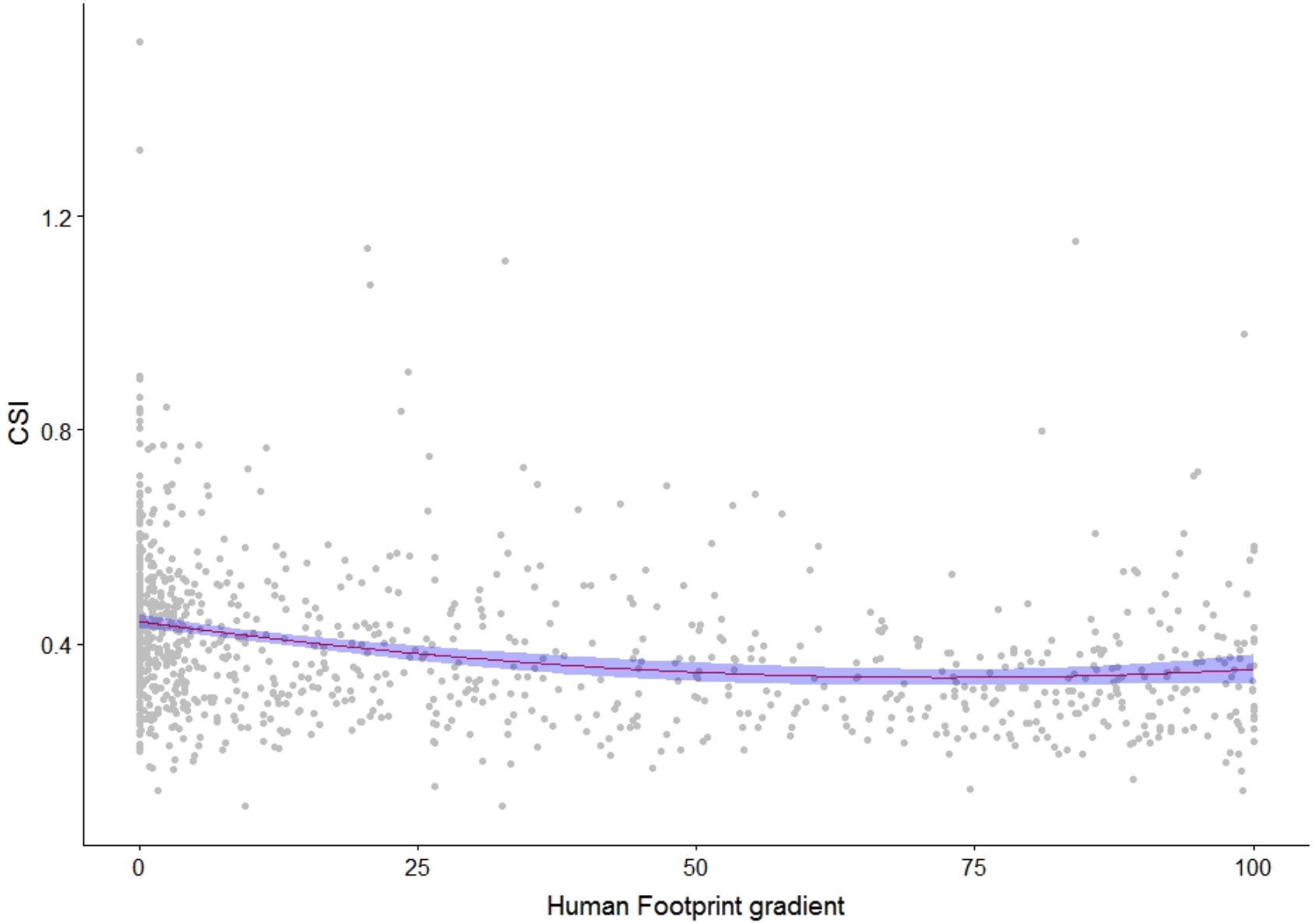
U-curve of community specialization is explained by the replacement of native species by invasive ones.

High correlation (r = 0.70) between HF % and Invasive species %



Calculation of CSI values and plot along the HF gradient - **Inverts**

- CSI = Community specialization index = mean of SSI present in each community



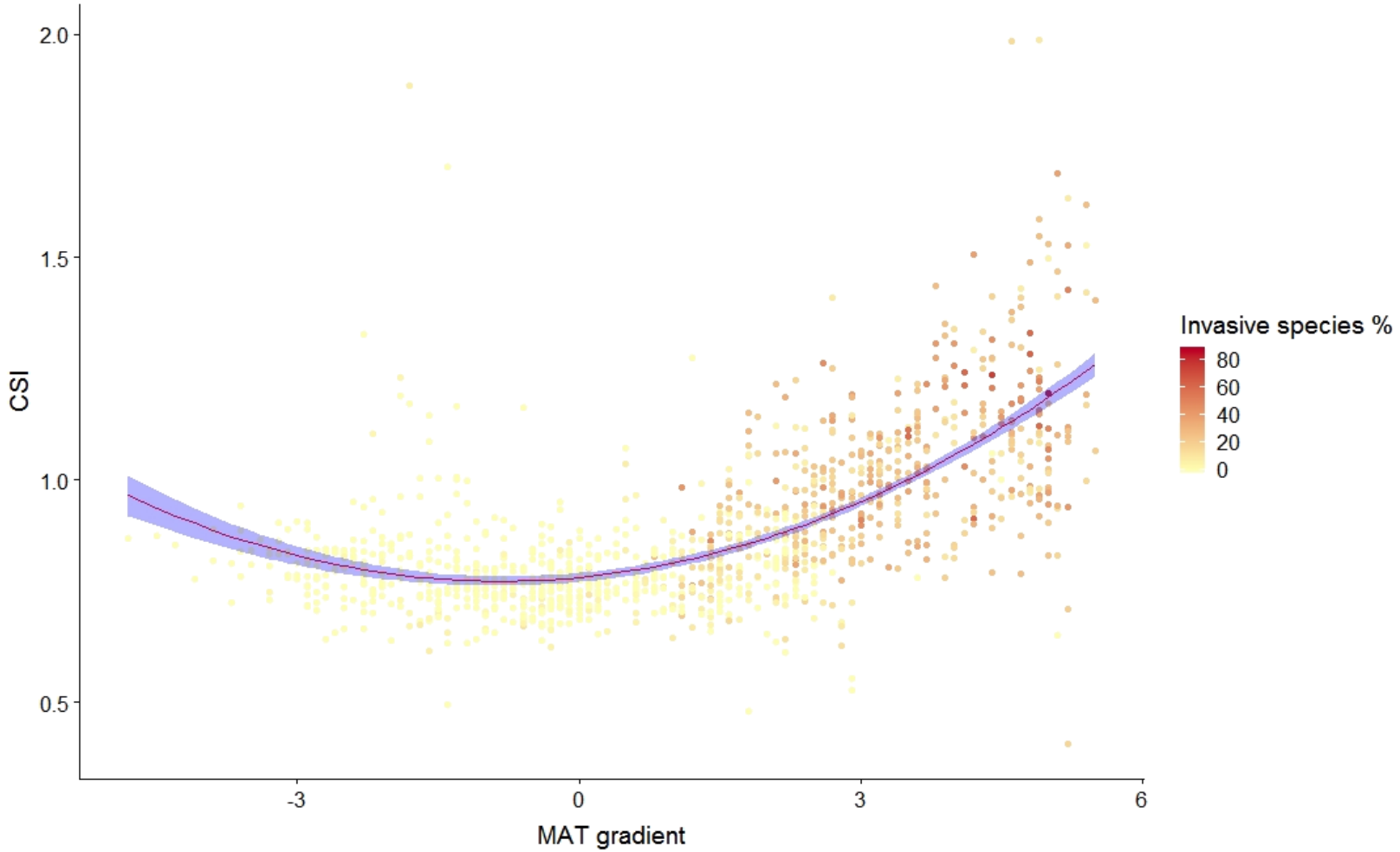
Interpretation:

U-curve pattern is low for invertebrates

Community specialization seem to decrease with gradient, no specialized species of disturbed part of the gradient.

Next steps ?

- Plants :
  - Look at how specialization of species is related to exotic status and preferences along the gradient, i.e. species with high specialization that preferred disturbed part of the gradient are really exotic ones?
  - Verify the found pattern and potential biases or confounding variable effect



Same pattern for MAT gradient?

(here only show wetlands protocol vegetation data because I did not have climate for the terrestrial sites (but could be the same values actually…)).