Demonstrating γ decreases with increasing β

Cape vs SWA publication

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Define a simple Jaccard distance function

```
jacc_dist <- function(a, b) {
  1 - (length(intersect(a, b)) / length(union(a, b)))
}</pre>
```

Create a species pool of richness n (analogous to the Cape or SWA)

```
n <- 1000
region <- 1:n
```

Define the range of richnesses m of the communities to draw from the pool (starting from stupidly a small 1 and ending at $\frac{1}{2}$ of regional richnes)

```
data \leftarrow data.frame(m = 1:(0.5 * n))
```

Draw communites a and b (a and b are analogous to smaller pixels, e.g. QDS), each of richness m, and calculate the richness γ of the larger scale community a+b (analogous to the larger scale pixel, e.g. HDS), and the turnover between a and b (i.e. the turnover "within" the larger scale community)

```
# Create empty columns to store results
data$richness_ab <- NA
data$jacc_dist_ab <- NA
for (m in data$m) {
  a <- sample(region, m)
  b <- sample(region, m)</pre>
  ab \leftarrow c(a, b)
  richness_ab <- length(unique(ab))</pre>
  jacc_dist_ab <- jacc_dist(a, b)</pre>
  # Store results in dataframe
  data$richness_ab[[m]] <- richness_ab</pre>
  data$jacc_dist_ab[[m]] <- jacc_dist_ab</pre>
  # Store adjusted results in dataframe too
  data$richness_ab_m_adjusted[[m]] <- richness_ab / m</pre>
  data$richness_ab_jacc_adjusted[[m]] <- richness_ab / jacc_dist_ab</pre>
}
```

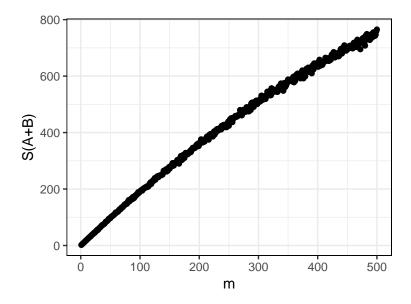


Figure 1: Plot of γ as a function of α

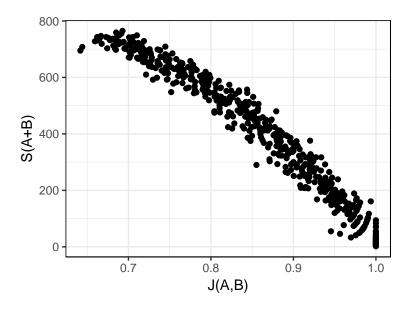


Figure 2: Plot of γ as a function of β

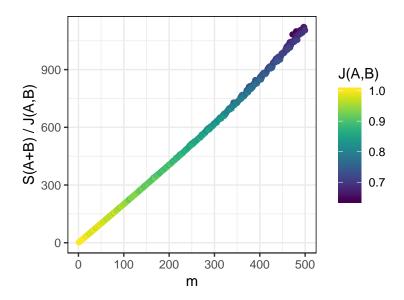


Figure 3: Plot of $\gamma_{adjby\beta}$ as a function of α

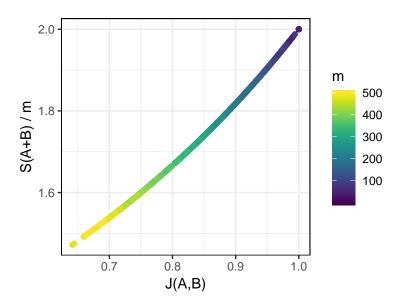


Figure 4: Plot of $\gamma_{adjby\alpha}$ as a function of β