**Order of scrips:**

1. CS\_00\_0a\_fish\_traits - Script for merging all traits values into a single csv file
2. CS\_00\_0b\_fish\_temporal\_sites - Script for merging biomass from temporal BRUV surveys
3. CS\_0a\_fish\_surveys - Script for preparing fish occurrence datasets
4. CS\_0b\_func\_space - Script for computing species positions in a multidimensional space according to their trait values
5. CS\_1a\_diversity\_spatial - Script for computing taxonomic and functional diversity and dissimilarity between habitats.
6. CS\_1b\_diversity\_temporal - Script for computing taxonomic and functional diversity and dissimilarity between years for kelp and no kelp sites
7. CS\_A\_figure1
8. CS\_B\_figure2
9. CS\_B2\_figure2\_sites
10. CS\_B3\_figure2\_turnover
11. CS\_C\_figure3

**Extras:**

1. 00\_trash – thermal affinity to add later
2. 0b\_functional\_space – old one, probably we won’t use this one
3. CS\_11a\_diversity\_spatial\_Hill - Script for computing taxonomic and functional diversity and dissimilarity between habitats using Hill approach