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Memòria del Treball Final deGrau

Effects of simulated herbivore pressure at different biogeographic levels on epifaunal communities associated with *Cymodocea nodosa*.

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Resum – Les praderies de fanerògames marines constitueixen un complex hàbitat costaner i juguen un paper clau en els ecosistemes marins a causa de la gran diversitat d'organismes que estan lligats a aquests ecosistemes. Actualment, molts d'aquests ecosistemes es troben en declivi i, en conseqüència, la biodiversitat que engloben. Les comunitats d'epifauna vàgils que viuen associats a les fulles de les praderies tenen un paper clau per transmetre la producció portada a terme per les herbes marines a nivells tròfics superiors. En aquest estudi, s'ha proposat examinar les variacions que poden arribar a partir les comunitats d'epifauna associades a les praderies de Cymodocea nodosa a causa de la pressió d'herbivoria simulada. Aquest experiment es va realitzar a diferents escales per obtenir diferències biogeogràfiques. Les localitzacions van ser Arinaga i Castillo de Romeral (a la regió de Gran Canària), Aucanada y Es Barcares (a la regió de Mallorca). El tractament d'herbivoria simulada va causar variacions en l'abundància tota de l'epifauna, la diversitat de l'epifauna i en l'abundància d'alguns grups taxonòmics. A més, les direccions de respostes dels grups taxonòmics no van ser uniformes entre localitats. Les biomasses de Cymodocea nodosa i els epífits semblen estar directament relacionades amb les comunitats d'epifauna.

Resumen - Las praderas de fanerógamas marinas constituyen un complejo hábitat costero y juegan un papel clave en los ecosistemas marinos debido a la gran diversidad de organismos que están ligados a estos ecosistemas. Actualmente, muchos de estos ecosistemas se encuentran en declive y, en consecuencia, la biodiversidad que abarcan. Las comunidades epifaunales vágiles que viven asociados a las hojas de las praderas tienen un papel crucial para transmitir la producción llevada a cabo por las hierbas marinas a niveles tróficos superiores. En este estudio, se ha propuesto examinar las variaciones que pueden llegar a sufrir las comunidades epifaunales associadas a praderas de Cymodocea nodosa a causa de la presión de herbivoría simulada. Este experimento se realizó a diferentes escalas para obtener diferencias biogeográficas. Las localizaciones fueron Arinaga y Castillo de Romeral (en la región de Gran Canaria), Aucanada y Es Barcares (en la región de Mallorca). El tratamiento de herbivoría simulada causó variaciones en la abundancia total de la epifauna, la diversidad de la epifauna y en la abundancia de algunos grupos taxonómicos. Además, las direcciones de respuestas de los grupos taxonómicos no fueron uniformes entre localidades. Las biomasas de Cymodocea nodosa y epifitos parecen estar directamente relacionadas con las comunidades de epifauna.

Abstract — Seagrass meadows constitute a complex coastal habitat and play a key role in marine ecosystems due to the high diversity of organisms that is linked to these ecosystems. Currently, many of these ecosystems are in decline and, consequently, their biodiversity is decreasing. Vague epifaunal communities that live associated with leaves of seagrass meadows have a crucial role in transmitting the production carried out by seagrasses to higher trophic levels. In this work, it has been proposed to study the variations that may occur to epifaunal communities associated with *Cymodocea nodosa* due to the pressure of simulated herbivorism. This experiment was carried out at different scales to obtain biogeographical differences. The locations were Arinaga and Castillo de Romeral (in the Gran Canaria region), Aucanada and Es Barcares (in the Mallorca region). The treatment of simulated herbivorism caused variations in the total abundance of epifauna, the diversity of epifauna and in the abundance of some taxonomic groups. In addition, the directions of responses of the taxonomic groups were not uniform among localities. Moreover, the biomasses of *Cymodocea nodosa* and epiphytes appear to be directly related to the epifauna communities.

1. Introduction

Human impacts have increased in recent times and, as a consequence, ecosystems have been affected. Marine ecosystems located close to the coast as coral reefs, mangrove forests, salt marshes and seagrass meadows support high impacts due to their geolocation and proximity to the impact sources (Balmford & Bond 2005, Orth et al. 2006, Beaumont et al. 2007). The impacts that these ecosystems receive are numerous such as nutrients and organic inputs, overfishing or susceptibility to climate change (seawater temperature rise or increased CO₂ concentration) (Jackson et al. 2007, Duarte 2002, Reynolds et al. 2018, Khan et al. 2016, Olsen et al. 2012). Mechanical damages (e.g. push nets or anchoring), aquaculture, eutrophication and salinity changes could produce changes in the meadow structures, erosion and finally a loss of the meadows (Duarte, 2002).

It is a well-known fact that seagrass meadows play a key role in marine ecosystems, due to the large number of ecological functions that they perform and services that they provide. In fact, seagrasses are considered "ecological engineers", as a result of physical, chemical and biological actions that seagrasses perform (Wright and Jones, 2006). Seagrasses influence physical factors such as the capture and stabilization of sediments and the attenuation of waves protecting coastal areas from turbulence (Nieuwenhuize & Hemminga 1990, Cabaço et al. 2010). The presence of seagrass meadows has an effect on the chemical factors in these ecosystems because they improve the quality of the water producing O₂ and incorporating nutrients into the water since they are responsible for the production of large amounts of organic carbon (Peduzzi & Vukovic, 2007). Furthermore, seagrass beds harbor a diverse community of small invertebrates, mainly formed by amphipods, copepods, annelids, gastropods, among other groups (Barberá-Cebrián et al. 2002, Tuya et al. 2006; Espino et al., 2011; Tuya et al. 2014b; Pnggonzalez, et al., 2014; Mateo Ramírez & García Raso, 2012). In some studies such as the one carried out by Lewis Allan W et al., (1983), a greater presence of macrofauna was observed in areas with seagrass meadows than areas without them. In fact, several studies claim that seagrass beds are a source of refuge for many organisms, some of them in danger of extinction (Duarte et al., 2008) and others with important economic value (Espino et al., 2011). Furthermore, some studies have shown that some fish

species have adapted their life cycle to the structural complex of seagrasses (Espino et al., 2011; Heck & Orth 2013).

The presence of epiphytic algae in the seagrass leaves is typically correlated with greater abundance and/or richness of species of the motile epifauna due to an increase in habitat complexity and a greater abundance of food (Nagle, 1968; Lewis and Hillingworth, 1982; Bell et al., 1984; Carr et al., 2011). Indeed, some organisms are strongly dependent on the epiphytes as their main food source (e.g. Tomas et al. 2006; Buza Jacobucci & Pereira Leite, 2014). In addition, it has been demonstrated the relevant role of mesograzers on seagrass meadows due to the crucial regulation that they perform as a result of their epiphytic consumption and, consequently, they control the competition for light and nutrients between seagrass and epiphytes (Neckles, et al., 1993). Epifaunal organisms are prey to a wide variety of consumers. In this way, they act as intermediate steps between primary and secondary producers transmitting the production carried out by seagrass meadow to higher levels in the trophic web. (Heck & Valentine 2006, Fernandez et al.; 2012). Moreover, seagrass leaves are also an important food source for macrograzers such as fish or sea urchins as well as megagrazers such as turtles, waterfowl and dugongs (Dugong dugong) (Moran & Bjorndal, 2005; Fourgurean et al., 2010).

Herbivory has fluctuations over time and space in different scales (Tomas et al., 2005; Prado et al. 2007,2008), but, even so, it has been shown that herbivory has important effects in seagrass meadows ecosystems. For instance, herbivory strongly influences the structure, composition and production of vegetation (McNaughton, 2001). Due to the alterations that herbivory can produce in these ecosystems, recent studies affirm that the effect of herbivory is a control factor in the fluctuations of dynamics population, composition, distribution and production, which can be stronger than temperature or salinity (Valentine & Heck 1999; Tomas et al., 2004; Gartner et al., 2013; Duffy et al. 2015; Carr et al. 2011). Therefore, herbivory effects can change the food and shelter resources that seagrass meadows offer to specific organisms that live associated with them and, consequently, alter epifauna abundances (McGlathery, 1995; Lal et al., 2010) as they.

Alter competition and predation, since habitat modification regulates the availability of food and shelter (Hixon & Menge 1991, Sirota & Hovel 2006, Gartner et al., 2013).

Cymodocea nodosa (Ucria) Ascherson is a dioecious seagrass distributed throughout the Mediterranean (including Balearic Islands), and adjacent eastern Atlantic coast but it is limited by southwestern Iberia, northwestern Africa and the Canary Islands and Maderia (Alberto et al., 2006; Mascaró et al., 2009; Tuya et al. 2014a). This species is considered a pioneer species colonizing soft bottoms under a wide range of environmental conditions. C. nodosa is the second species of seagrasses in terms of abundance in the Mediterranean Sea (being Posidonia oceanica the first one; Scipione & Zupo, 2010) and the most important macrophyte in the Canary Island. (Reyes et al.,1995; Barbera et al. 2005). The evolutionary origin of this seagrass was the Indian Ocean (Red Sea), by divergence, because of its proximity, colonized the eastern Mediterranean, and spread across the Atlantic to the west, thus constituting the central and peripheral part of this seagrass (Masucci & Arnaud-haond 2012, Alberto et al. 2006). As previously mentioned, many studies have shown that herbivory has effects linked to the performance of seagrass and its ecology, as it also happens in the case of C. nodosa (Cebrián et al. 1996, Fernandez et al. 2012, Tuya et al., 2013). Even so, the effects of herbivory on the epifauna associated with C. nodosa have not been studied yet and remain unknown.

The aim of this study is to evaluate the effects of simulated herbivory and biogeographic location on epifaunal organisms associated with the seagrass *C. nodosa*. To examine these effects, meadows locations have been selected in two distinc regions within the distribution area of *C. nodosa*.

2. Methods

2.1 Study sites

Four *C. nodosa* meadows from two regions (Canarias and Mallorca) were selected in order to study the effects of simulated herbivore pressure at different biogeographical scales. The meadows from Canary Island belong to the peripheral zone of distribution of *C. nodosa*, whereas the meadows located in Baleares are within the central part of the distribution range of *C. nodosa* (Alberto et al., 2006; Masucci & Arnaud-haond, 2012).

In Canary Island, the two meadows selected were located in Gran Canaria Island. The first sampling site was Arinaga (27º51'N 15º24'W), where there is a meadow of *C. nodosa* with an area of 1.5ha, located at 10m depth and with a shoot density of 424±83 m². The second sampling sites was Castillo del Romeral (27º47'N 15º29'W), where a meadow with an area of 25ha, located at 5m depth and with a shoot density of 787±156 m² is found. These two location are situated in the southeast region of Canary Island (Figure 1, Canary Island map). The other two meadows are located in the northern region of Balearic Island.



Figure 1. Location the C. nodosa meadows in Gran Canaria: Arinaga (green) and Castido del

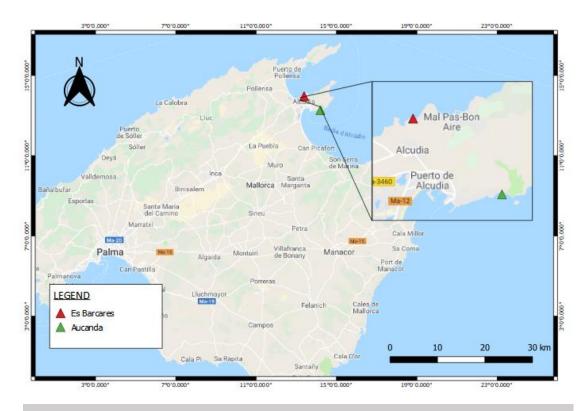


Figure 2. Location the C. nodosa meadows in of Mallorca: Es Barcares (red) and Aucanda (green).

One of them was located in Aucanada (39º50'N 3º10'E), where the meadow present an area of 2.2ha, located at 2.7m and with a shoot density of 1183±204 m². The last study meadows was located in Es Barcares (39º51'N 3º07'E), where the meadow has an area of 1.6ha, located at 3.5m and with a shoot density of 756±188 m² is found (Tuya et al., 2018) (Figure 2, Balearic Island map).

2.2 Field sampling methods

Different levels of herbivory were simulated in each *C. nodosa* meadows though a manipulate experiment. This experiment and the subsequent sampling were performed by SCUBA diving. First of all, 10 plots (1 x 1 m) were put in the seabed haphazard, separated from each other by 2 meters. Then, high herbivory level was set up in 5 plots cutting the seagrass leaves at 80%, whereas the rest of plots were established as control (i.e. natural low herbivory level). The experiment was carried out during the summer season, from June to September of 2016, coinciding with highest activity of fish grazing (Tomas et al., 2005) and leaves were cut once a month to maintain treatment levels. At

the end of the experimental period, a 0.04m² quadrat with a cotton bag mesh (size 250µm) was placed in the center of each plot (avoiding any border effects) to collect the epifauna. Associated fauna and aboveground biomass were caught inside the mesh bag by carefully, cutting the seagrass leaves just above the sediment surface with scissors (Figure 3). Collected samples were kept in individual plastic bags filled with seawater and transported to the laboratory in coolers for further sorting.



Figure 3. Sampling of epifauna after having applied the simulated herbivorous treatment.

2.3 Laboratory analyses

All plastic bags with the corresponding samples were stored in the freezer until further manipulation in the laboratory. The content of each bag including leaves of *C. nodosa* were rinsed with freshwater using a three-rinse cycle separating epifauna from leaves (Carr et al., 2011). Then, samples were filtered through a 50 and 125 micrometers mesh sieve to separate epifauna by size. The different samples resulting from the separation by size were preserved in 70% ethanol. Carefully, epiphytes were scrapped off the leaves by a glass slide and were placed in a previously weighed filter. Epiphytes and leaf biomass were dried in an over for 48 hours at 60°C, and subsequently weighted (Dry Weight, DW). Epifauna specimens with a size of 500 micrometers or larger were sorted into taxonomic units and their abundance was quantified using a binocular microscope (Tuya et al., 2013)(Figure 4).



Figure 4. Laboratory analysis process. Rinse the leaves of *C. nodosa* with fresh water causing an osmotic shoc and the subsequent separation of the epifauna from the leaves (1). Separation of epyphites and leaves (2). Filtration through a tapestry of 500 micrometers of the epifauna (3). Classification of taxonomic groups by binocular microscope (4).

2.4 Statistical analyses

In order to examine the existence of significant differences between biomass of C. nodosa (gDW/100cm²) and biomass of epiphytes (gDW/100cm²) across the factors region, location and treatment, three-way multivariate permutational ANOVA (PERMANOVA) was carried out. The correlation between the biomass of epiphytes, the biomass of C. nodosa, the total abundance of individuals (ind/100cm²) as well as the taxonomic richness (species/100cm²) was analyzed by the Spearman correlation test. Parametric analyses were used when normality (Shapiro Wilk's and Q-Q norm quantiles based on residuals (Annex S1; Annex S3; Annex S5)) and homoscedasticity (Levenne test) of the data were detected. Then, a three-way Analysis of Variance (ANOVA) was performed to compare the differences in total abundance of individuals (ind/100cm²), taxa richness (species/100cm²), and different diversity measures such as Pielow eveness (J) (Feldman, 2006) and Shannon Wiener diversity index (H') (Shannon and Weaver, 1963) across the factors of Region (fixed factor; 2 levels), Locations (nested within region; 2 levels) and clipping treatment (fixed factor; 2 levels). Fourth root transformation data were used to generate a similarity matrix based on Bray Curtis index. Subsequently, multidimensional scaling (MDS) representation and three-way multivariate permutational ANOVA (PERMANOVA) were performed to examine the differences in the structure of the epifauna community associated with the different C. nodosa meadows. The design was the same than ANOVA analysis. To conclude, a similitude test (SIMPER routine) was used to determine which epifauna groups contributed to a greater differences per regions and between them. Two-way Analysis of Variance (ANOVA) was performed to compare the differences in the taxonomic groups that produce a greater differences between region (fixed factor, 2 levels) and clipping treatment (fixed factor, 2 levels). When the criteria of normality and homoscedasticity were not respected Kruskal-Wallis test was used.

Univariate statistical analyses, homogeneity test and homoscedasticity test were performed using Rstudio software package (Version 1.1.463). Graphics and maps were plotted with RStudio, Excel and QGIS 3.6. Diversity index, multivariate methods (PERMANOVA, MDS), and SIMPER analyses, were performed with the PRIMER 7 software package.

3. Results

3.1 Cymodocea nodosa and epiphytes biomass

Biomass of *Cymodocea nodosa* presents significant differences between Region, Location and Treatment factors (Table 1). Higher abundance of seagrass is found in Gran Canaria, is also the region where the treatment factor affects with greater magnitude (Figure 5). Epiphyte biomass presents differences between region, location and treatment, according to the statistical results made with three-way multivariate permutational ANOVA (PERMANOVA) (Figure 6, Table 1).

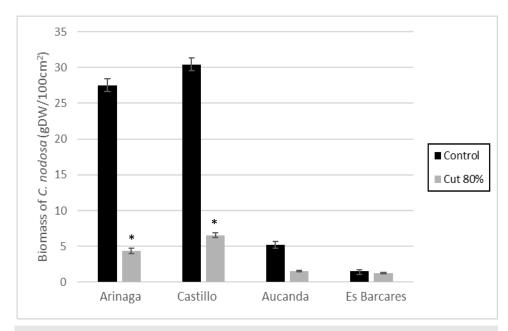


Figure 5. Biomass of *Cymodocea nodosa* (g*100cm $^{-2}$) (mean and SE) at the four location Arinaga, Castillo, Aucanada i Es Barcares and across clipping treatments (Control and Cut (80%)). Data (n= 18 plots per treatment). Significant values of Wilcoxon rank sum test (*).

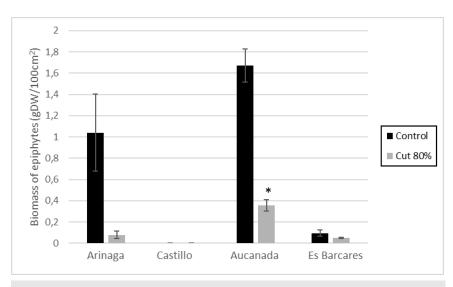


Figure 6. Biomass of epiphytes (g*100cm⁻²) (mean and SE) at the four location Arinaga, Castillo, Aucanada i Es Barcares and across clipping treatments (Control and Cut (80%)). Data (n= 18 plots per treatment). Significant values of Wilcoxon rank sum test (*).

Table 1. Results of three-way multivariate permutational ANOVA (PERMANOVA) examining differences in biomass of *C. nodosa* (gDW/100cm²) and biomass of epiphytes (gDW/100cm²) between Region (Re), locations between regions (Loc(Re)), treatment (Tr) and their interactions. Both variables have need a pre-treatment (square root). Pseudo-F = pseudo-F values, df = degrees of freedom, MS = mean squares, unique perms = numbers of permutations. Significant p-values are highlighted in bold.

Source of variation	df	MS	Pseudo-F	P (perm)	Unique perms
Biomass of C. nodosa					
Re	1	0.5033	239.85	0.0001	9840
Tr	1	0.2297	109.49	0.0001	9850
Loc(Re)	2	0.0171	8.1255	0.0017	9951
RexTr	1	0.0636	30.298	0.0001	9848
Loc(Re)xTr	2	0.0018	0.8672	0.4205	9954
Residuals	28	0.0021			
Total	35				
Biomass of epiphytes					
Re	1	0.31423	9.7281	0.0042	9831
Tr	1	0.13875	4.2956	0.0347	9849
Loc(Re)	2	0.27206	8.4224	0.0015	9963
RexTr	1	0.00807	0.2499	0.6589	9874
Loc(Re)xTr	2	0.05287	1.6369	0.21	9956
Residuals	28	0.03230			
Total	35				

The results obtained from the Spearman correlation test determine that the abundance of individuals (N) and the taxonomic richness (T) are positively correlated with epiphytes biomass, whereas, the results between the abundance of epiphytes and the abundance of the seagrass show a significant negative correlation (Table 2, Figure 7, Figure 8)

Table 2. Results of the Spearman correlation test on the relationship between biomass of epiphytes (gDW/100cm²) and biomass of *Cymodocea nodosa* (g/100cm²), total abundance of individuals (N; Ind/100cm²) and Taxon Richness (T; species/100cm²). S = Spearman correlation rank, rho = Spearman coefficient rank. P-value in bold indicates positive correlation.

Source of variation	S	p-value	rho	
Epiphytes				
Leaf	10509	0.035	-0.352472	
N	4406.7	0.008367	0.4328629	
Т	2817.6	0.00002909	0.6373765	

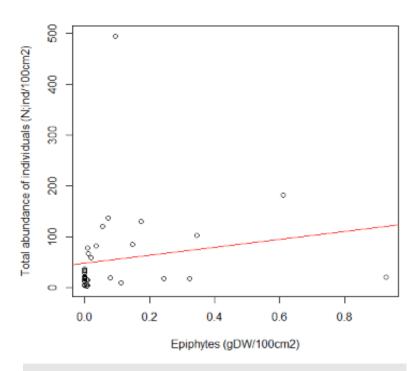


Figure 7. Relationship between Total abundance of individual (N; $ind/100cm^2$) and Epiphytes biomass (gDW/100cm⁻²). R²=0.0299

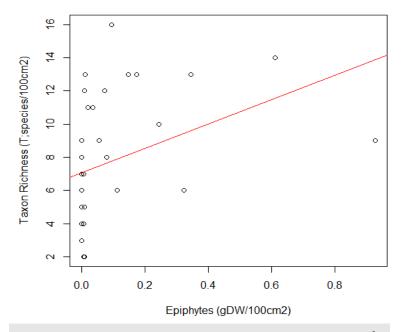


Figure 8. Relationship between Taxon Richness (T; species/100cm²) Epiphytes biomass (gDW/100cm⁻²). R²=0.1424

3.2 Epifaunal groups

A total of 1952 individuals were counted and classified in taxonomic groups resulting in 543 Nematoda, 305 Polychaeta, 250 Copepoda, 220 Prosobranchia, 195 Gammaridea, 152 Caprellidea, 70 Cumacea, 55 Tanaidacea, 41 Isopoda, 31 Bivalvia, 22 Ostracoda, 13 Pycnogonida, 13 Arachnida, 11 Echinoidea, 8 Sipunculidea, 6 Sagittoidea, 5 Anomura, 5 Caridea, 3 Mysidacea, 3 Brachyura, 1 Clitellata (Hirudinea).

The abundance of individuals (N; ind/100cm²) presents significant differences between region and location. In addition, the effects of the treatment factor (clipping) were different between regions (i.e. nearly significant interaction treatment x region. Whereas abundance of individuals tended to decrease under clipping, in location of Aucanda, in Mallorca, the treatment cut 80% exhibited higher abundance of individuals than the control (Table 3; Figure 9 Post Hoc; p>0.05 Annex S4).

Table 3. Results of univariate Three-way nested ANOVAs comparing differences in total abundance of individuals (N; ind/ $100cm^2$), taxa richness (T; species/ $100cm^2$), Pielow and Shannon Wiener (H') per region (Re), location (Loc) and treatment (Tr) and their interactions. Region factor (Re) is nested within location (Loc(Re)). df = degrees of freedom, SS = sum of squares. Significant p-values are highlighted in bold.

Source of variation	n	df	SS	F	p-value
Abundance (N,ind/100 cm²)	36				
Re		1	1.667	9.330	0.00491
Loc(Re)		2	1.992	5.574	0.00917
Tr		1	0.128	0.716	0.40453
Re:Tr		1	0.069	0.388	0.53859
Loc (Re):Tr		2	1.140	3.190	0.05647
Residuals		28	5.004		
Taxon Richness (T)	36				
Re		1	124.69	13.020	0.00119
Loc(Re)		2	81.61	4.261	0.02424
Tr		1	0.40	0.042	0.83927
Re:Tr		1	2.81	0.294	0.59216
Loc (Re):Tr		2	15.08	0.787	0.46487
Residuals		28	268.15		
Pielou's eveness (J)	36				
Re		1	0.00901	2.567	0.12034
Loc(Re)		2	0.04148	5.912	0.00721
Tr		1	0.00055	0.156	0.69610
Re:Tr		1	0.02063	5.879	0.02202
Loc (Re):Tr		2	0.03391	4.834	0.01573
Residuals		28	0.09823		
Shannon Wiener (H')	36				
Re		1	0.335	2.256	0.1443
Loc(Re)		2	0.303	1.022	0.3730
Tr		1	0.088	0.590	0.4490
Re:Tr		1	0.791	5.332	0.0285
Loc (Re):Tr		2	0.057	0.193	0.8256
Residuals		28	4.156		

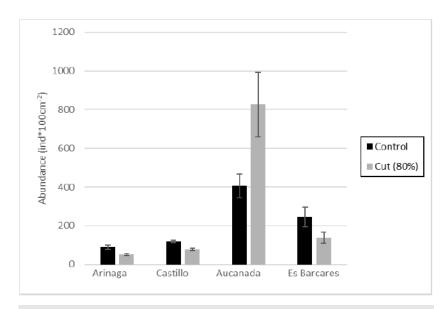


Figure 9. Total abundance of individuals (N; ind*100cm⁻²) (mean and SE) at the four Location and across clipping treatments (Control and Cut (80%)). Data (n= 5 plots per treatment).

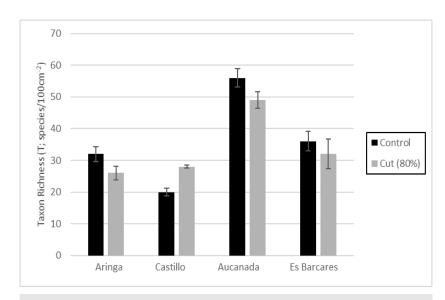


Figure 10. Taxon Richness (T; species/100cm²) (mean and SE) at the four Location and across clipping treatments (Control and Cut (80%)). Data (n= 5 plots per treatment).

The number of taxa (T/species/100cm²) showed significant differences between region, with Mallorca presenting higher taxon richness values than Gran Canaria (Table 3; Figure 10; Post Hoc; p>0.05 Annex S4). In the case of Shannon Wiener index the location of Aucanada shows the highest value of this index. The location of Castillo de Romeral shows a smaller value of Shannon Wiener index and it was the unique location where

the effect of the treatment cut 80% showed higher values than control (Table 3; Figure 11; Post Hoc; p>0.05 Annex S4). The effects of the 80% clipping treatment in Pielow's evenness index varies by Region (Table 3; Figure 12; Post Hoc; p>0.05 Annex S4), where there were also significant differences between regions.

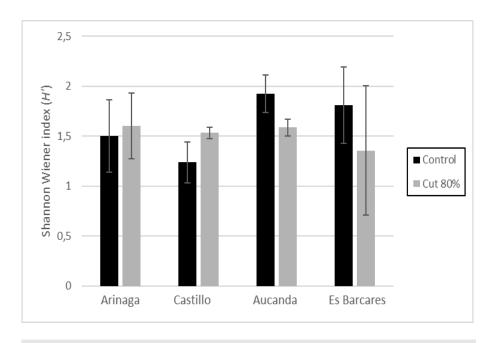


Figure 11. Shannon Wiener index (H') (mean and SE) at the four Location and across clipping treatments (Control and Cut (80%)). Data (n= 18 plots per treatment).

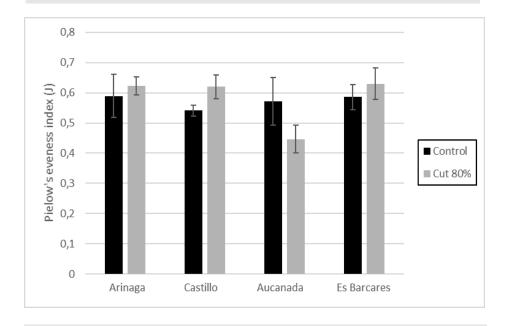


Figure 12. Pielow's eveness index (J) (mean and SE) at the four Location and across clipping treatments (Control and Cut (80%)). Data (n= 18 plots per treatment).

Pielow's eveness index (*J*) showed significant values in terms of the locality factor between Aucanda and Es Barcares, and the interaction between locality (region factor nested within) and treatment between Aucanda and Es Barcares after 80% clipping treatment (Significant results of the post hoc analysis of Tukey HSD test). The values of Pielow's eveness are greater in the treatment of 80% clipping with the exception of Aucanada where Pielow's evenness index is greater in the control treatment.

The structure of the community displayed significant differences between regions but not between treatment levels (Table 4). The MDS show a coherent grouping arrangement according to the regions (Figure 13).

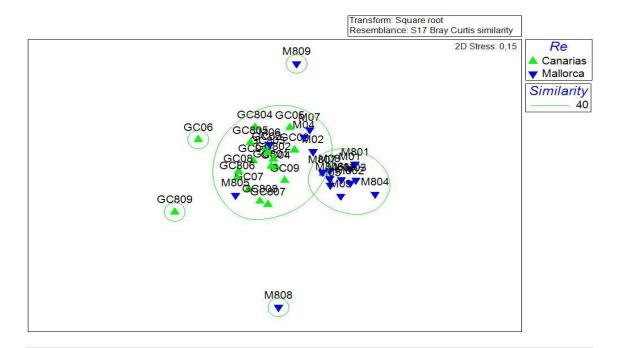


Figure 13. Multidimensional scaling plot from the transformation of the data to the fourth root and the generation of the similarity matrix generated by an index of similarity of Bray Curtis considering the abundance of the different groups of epifauna at each site according to the region (green=Canarias, blue=Mallorca) and treatment factors (Control, Cut 80%). Circles group samples under with a 40% similarity array. Data (n= 5 plots per treatment).

Table 4. Results of three-way multivariate permutational ANOVA (PERMANOVA) examining differences in the abundance of each taxon (ind/ $100cm^2$) between Region (Re), locations between regions (Loc(Re)), treatment (Tr) and their interactions. Pseudo-F = pseudo-F values, df = degrees of freedom, MS = mean squares, unique perms = numbers of permutations. Significant p-values are highlighted in bold.

df	MS	Pseudo-F	P(perm)	Unique perms
1	14373	10.614	0.0001	9947
2	4836.1	3.5715	0.0001	9909
1	1880.1	1.3885	0.2017	9941
1	833.07	0.61523	0.7762	9939
2	1185.7	0.87561	0.6037	9922
28	1354.1			
35	66631			
	1 2 1 1 2 28	1 14373 2 4836.1 1 1880.1 1 833.07 2 1185.7 28 1354.1	1 14373 10.614 2 4836.1 3.5715 1 1880.1 1.3885 1 833.07 0.61523 2 1185.7 0.87561 28 1354.1	1 14373 10.614 0.0001 2 4836.1 3.5715 0.0001 1 1880.1 1.3885 0.2017 1 833.07 0.61523 0.7762 2 1185.7 0.87561 0.6037 28 1354.1

Nematoda, Caprellidea, Polychaeta, Copepoda, Prosobranchia, Gammaridea, Cumacea and Tanaidacea were responsible of the 70% differences between Gran Canaria and Mallorca. Therefore, the differences found between regions depend mainly on a group of determined taxa (Figure 5). Furthermore, the results of three-way multivariate permutational ANOVA (PERMANOVA) that there is no effect of clipping on the community structure overall, although, there were some specific taxonomic groups in which the treatment factor has effects on their abundances (Figure 14).

For instance, the Nematoda taxon presents significant differences at the level of both the region and treatment factors (Table 6; Figure 14; Post Hoc analysis p-value<0.05 Annex S6).

Table 5. Results of SIMPER test for similarity between regions: Gran Canarias (GC) and Mallorca (M). Av. Abund = abundance of each unit taxon, Av. Diss = average dissimilarity, Diss/SD = quotient of dissimilarity and standard deviation, Contrib. % = contribution of each taxon to the overall dissimilarity between high and low groups, Cum% = cumulative contribution.

Group Gran Canarias & Mallorca Average of dissimilarity = 64.93							
	Group GC	Group M					
Species	Av. Abund	Av. Abund	Av. Diss	Diss/SD	Contrib. %	Cum. %	
Nematoda	0.06	3.93	9.72	1.38	14.98	14.98	
Caprellidea	0.44	2.39	6.66	1.55	10.25	25.23	
Polychaeta	0.39	2.67	6.64	1.13	10.23	35.46	
Copepoda	1.32	2.70	6.25	1.34	9.62	45.08	
Prosobranchia	2.12	2.19	5.90	0.95	9.09	54.17	
Gammaridea	2.25	1.89	5.28	0.89	8.13	62.30	
Cumacea	0.00	1.47	3.86	1.11	5.94	68.24	
Tanaidacea	0.40	1.28	3.81	1.21	5.87	74.11	
Bivalvia	0.00	1.01	3.01	1.07	4.63	78.74	
Isopoda	0.65	0.90	3.00	0.96	4.61	83.35	
Echinoidea	0.47	0.00	1.93	0.62	2.98	86.33	
Ostracoda	0.22	0.61	1.83	0.85	2.82	89.15	
Arachnida	0.32	0.30	1.51	0.69	2.32	91.48	
Pycnogonida	0.08	0.39	1.18	0.60	1.81	93.29	
Caridea	0.25	0.00	1.13	0.42	1.73	95.02	
Sipunculidea	0.00	0.34	0.80	0.60	1.23	96.25	
Mysidacea	0.06	0.11	0.60	0.31	0.93	97.18	
Sagiitoidea	0.00	0.27	0.59	0.50	0.91	98.09	
Anomura	0.13	0.08	0.57	0.35	0.87	98.97	
Brachyura	0.06	0.11	0.53	0.39	0.82	99.79	
Hirudinea	0.00	0.06	0.14	0.24	0.22	100.00	

In addition, the taxonomic group Prosobranchia displayed significant differences only at the level of treatment factor, being less abundant under clipping (Table 7; Figure 14; Post Hoc analysis p-value<0.05 Annex S7).

On the other hand, the taxonomic group Gammaridea did not show significant differences in any factor involved. On the other hand, the majority of taxonomic groups such as Caprellidea, Cumacea, Copepoda and Tanidacea presented significant differences at the level of region factor, generally being more abundant in Mallorca than in Gran Canaria (Table 6; Table 7).

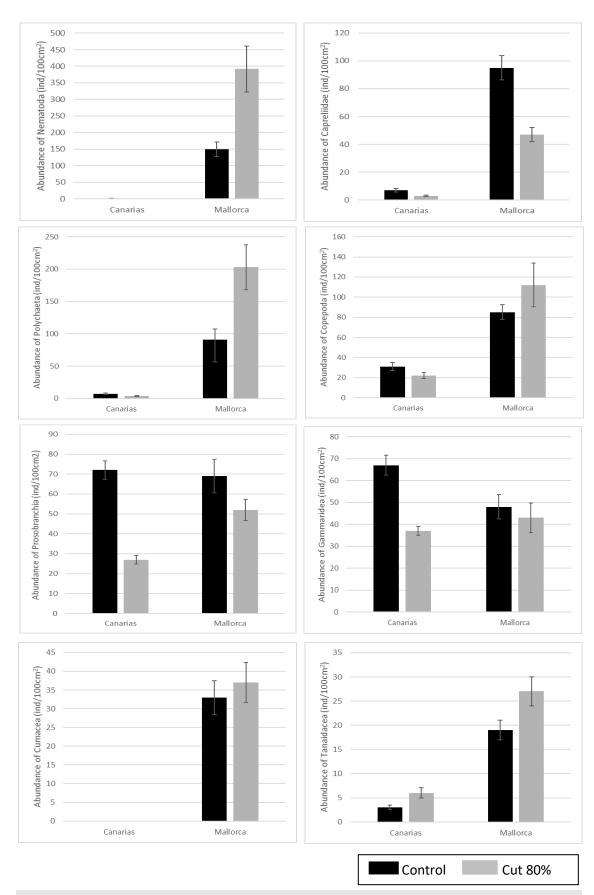


Figure 14. Results of Abundance of individuals (ind/100 cm²) of the Nematoda, Caprellidea, Polychaeta, Gammaridea, Copepoda, Prosobranchia, Gammaridea, Cumacea, Tanaidacea taxon groups found in *Cymodocea nodosa* across regions and experimental treatments.

Table 6. Results of the non-parametric Kruskal-Wallis H test on the Nematoda, Polychaeta, Caprellidea, Cumacea and Tanaidacea abundance (ind/100cm²) across the fixed factors region (Re) and clipping reatment (Tr). df = degrees of freedom, $\chi 2$ =Kruskal-Wallis Chi-squared. Significant p-values are highlighted in bold.

Source of variation	df	χ^2	p-value
Nematoda			
Re	1	19.202	0.000011176
Tr	1	3.8467	0.04984
Polychaeta			
Re	1	7.3547	0.006689
Tr	1	0.04979	0.8234
Caprellidea			
Re	1	16.032	0.00006227
Tr	1	1.6669	0.1967
Cumacea			
Re	1	16.594	0.00004629
Tr	1	0.00035567	0.985
Tanaidacea			
Re	1	8.2606	0.004052
Tr	1	0.36612	0.5451

Table 7. Results of univariate Three-way nested ANOVAs comparing differences in Prosobranchia (pretreatment Ln+1), Gammaridea (pre-treatment Ln+1) and Copepoda (pre-treatment Ln+1) abundance (ind/ $100cm^2$) per region (Re), location (Loc) and treatment (Tr) and their interactions. Region factor (Re) is nested within location (Loc(Re)). df = degrees of freedom, SS = sum of squares. Significant p-values are highlighted in bold.

Source of variation	n	df	SS	F	p-value
Prosobranchia	36				
Re		1	0.000	0.001	0.9795
Loc(Re)		2	0.896	0.669	0.5201
Tr		1	3.097	4.628	0.0402
Re:Tr		1	0.783	1.170	0.2887
Loc (Re):Tr		2	2.788	2.084	0.1434
Residuals		28	18.736		
Gammaridea	36				
Re		1	0.869	1.505	0.230
Loc(Re)		2	1.224	1.060	0.360
Tr		1	0.920	1.593	0.217
Re:Tr		1	0.074	0.129	0.722
Loc (Re):Tr		2	1.825	1.580	0.224
Residuals		28	16.170		
Copepoda	36				
Re		1	7.292	12.889	0.001246
Loc(Re)		2	14.484	12.801	0.000113
Tr		1	0.625	1.104	0.302306
Re:Tr		1	0.061	0.109	0.744303
Loc (Re):Tr		2	1.659	1.466	0.247952
Residuals		28	15.841		

4. Discussion

The results obtained in this study suggest that the factor that regulates the structure and composition of the epifauna communities is the different biogeographical regions. Moreover, our results also suggest that the clipping treatment mimicking the herbivory pressure modifies the total abundance and diversity index of epifauna. Unexpectedly, the effect of the clipping treatment has had opposite outcomes depending on the location (Table 3). For instance, the abundance of the Prosobranchia taxonomic group decreased upon clipping treatment whereas, the abundance of Nematoda taxonomic group increased (Table 7). These data support the idea that epifauna is very sensitive to environmental variations as it was previously shown in different studies, such as the study carried out by Tanner et al. (2006). This study unveiled that the variation of the epifauna composition is affected by the patch of meadows, landscape characteristics and wave exposure variable. Our study suggests a new factor to the epifauna community regulation: the herbivory pressure.

However, the differences found in this study could be caused by the location of the analyzed regions. Firstly, because these regions are located in archipelagoes but with the difference that Mallorca is located very close to the Iberian Peninsula. The study carried out by Benedetti-Cecchi et al. (2003) showed that both the algae and the set of invertebrates differed from the islands to the continental shallow waters, since isolated islands provide fundamentally different habitat patterns of those of the continental shallow waters.

Secondly, these locations are exposed to different hydrodynamism. For instance, the open sea location of the *Cymodocea nodosa* meadow in Gran Canaria (southeast region) causes the fragmentation of seagrass probably due to hydrodynamic effects caused by waves and tides. These damaging hydrodynamic effects may be stronger in Gran Canaria than in Mallorca because *Cymodocea nodosa* meadows located in Mallorca are in areas more protected from hydrodynamism such as caves and bays (Tuya et al., 2014b). This weaker hydrodynamism may explain the higher abundance of individuals in Mallorca compared to Gran Canaria (Figure 9, Table 3).

Our data also suggest a positive effect of biomass of *C. nodosa* and epiphytes in the abundance and taxa richness of epifaunal communities (Figure 7, Table 2). Indeed, several studies have previously shown that changes in the habitat structure (epiphytic biomass and leaf biomass) affects the diversity and abundance of the epifauna community (Gartner et al. 2013; Edgar and Robertson, 1992; Heck and Wetstone, 1977; Stoner, 1980; Attrill et al. 2000). Specifically, some studies link epifaunal composition to the biomass of epiphytes instead of leaf biomass because epiphytes provide an increased complexity to habitat that promotes epifauna diversity (increase in shelter and food) (Nagle, 1968; Novak, 1982; Lewis & Hollingworth, 1982; Bell et al., 1984). Interestingly, our data show differences between the leaf biomass and epiphytes biomass between regions and locations (Table 1). These results suggest that there are latitudinal differences in the growth of alga as have been previously demonstrated in others studies (Heck & Valentine, 2006; Gartner et al. 2013).

The abundance of epifaunal individuals was significantly influenced by the clipping treatment, but the responses differed amongst localities (Table 3). Whereas clipping tended to decrease the total abundance of epifauna, in Aucanada (in the Mallorca region), total abundance was not only the highest of all localities, but was also where clipping actually increased total abundance. In fact, the clipped plots of Aucanada exhibited the highest overall abundance recorded. Similarly to abundance, response of epifaunal diversity patterns to the clipping treatment also differed between regions. The Shannon Wiener index (H') before the treatment tended to be higher in the Mallorca sites compared to Gran Canaria (although they were still relatively low specific biodiversity values; H' < 2). In these Mallorca locations, the effects of clipping tended to decrease diversity. However, the Shannon Wiener index tended to increase in the clipping 80% treatment in the localities of Gran Canaria (Table 3). This unexpected increase in the Shannon Wiener index could be explain by the structural complexity of macroalgal beds, since the excessive vegetative development of algae could prevent the access of new individuals due to the high compression of algal ramifications. This increase in the Shannon Wiener index is not observed in Mallorca due to the initial conditions of lower biomass and lack of excessive development of C. nodosa. The increase in epifauna diversity caused by the lower algal development was also observed in cold months when the structural complexity of macroalgal beds could facilitate the increase of the number of individuals (Mateo-Ramírez et al., 2018).

Regarding Pielow's evenness (*J*), higher values were observed under the 80% clipping treatment except in the case of Aucanada. This result suggest that the removal of leaf biomass produces a modification on habitat structure. This restructuring effect affects negatively the abundance of dominant taxa such as Prosobranchia (Table 3). This decrease in Prosobranchia abundance after the 80% clipping treatment would allow the increase in the abundance of other taxonomic groups that have lower values before clipping treatment, such as Isopoda, Copepoda, Tanaidacea and Caprelidae. Similarly to Prosobranchia, other dominant taxa, such as Caprellidae or Gammaridea, showed a decreased abundance after the 80% clipping treatment. However, significant differences were only observed in the Prosobranchia taxon. On the other hand, as expected, the several taxa, such as Nematoda, Polychaeta, Copepoda, Cumacea and Tanaidacea showed higher abundance values after the 80% clipping treatment compared to the control. Therefore, these changes in the abundances of the different taxa would explain the increase in the value of Pielow's eveness index after clipping treatment, since after clipping treatment the different taxa populations tend to equal.

There are different reasons to explain the population changes of the different taxa after clipping treatment. For instance, Prosobranchia population decreases because this taxon population is highly linked to the abundance of seagrass (Urra et al., 2013a; Marina et al., 2012; Urra et al., 2013b). On the other hand, Nematoda population increases probably due to the decrease of predatory pressure on this taxon by fishes and other predators that require high abundance of seagrass in their habitat to subsist (Mateo-Ramírez et al., 2018).

Unexpectedly, the Pielow's eveness index in Aucanada location decreases after clipping treatment. This result is apparently contradictory to the rest of the data obtained in other locations. However, Aucanada is different from the rest of locations before the clipping treatment, since Aucanada shows a higher biomass of epiphytes and a higher abundance of epifauna than the rest, even with a lower seagrass biomass. The Aucanada data obtained before the clipping treatment suggest that Aucanada is a specially rich habitat probably due to the high deposition of organic material of a river

that flows into the bay less than 500 meters away from the study area. The clipping treatment would facilitate the access to this organic material to the epifauna. Therefore, and in contrast to the rest of locations, the clipping treatment in Aucanada increases the abundance of almost all the taxa studied. This unexpected increase in the abundance of epifauna after clipping treatment would explain the decrease in the Pielow's eveness index in Aucanada.

The structure and complexity of seagrass habitat is defined by two main group of factors: the natural factors, sucs as surface area, number of leaves per seagrass shoot, patch mosaic, leaf growth taxa, leaf biomass, latitude or temperature (Sirota & Hovel, 2006; Larkum et al., 2006; Tanner, 2006); and the anthropic factors, such as organic matter or pollution (Duarte, 2002). The results obtained during this study suggest that the effect of herbivorism is a factor involved in a broader and more important way in the seagrass habitat complexity regulation. The data of this study suggests that the herbivorism could be regulating the abundance and composition of the epifauna community in the seagrass meadow. However, the degree of influence that the herbivorism has on abundance of epifauna, taxonomic richness or structure of the community remains still unclear. Further investigations should clarify the role of herbivorism in the regulation of the epifauna community that live on the leaves of seagrass. The importance of the epifauna in the proper regulation of the seagrass ecosystem is widely known and that is why it is important to continue studying the different factors that affect epifauna in order to assure the conservation of these important marine habitats.

5. Conclusions

With the results obtained in this study, it has been possible to demonstrate that the pressure of herbivory can cause impacts on the mobile fauna that lives above the leaves of *C. nodosa* at a regional level and, specifically, only in some taxonomic groups. The results have shown differences in the total abundance and taxa richness between the regions of Mallorca and Gran Canaria. In addition, the abundance of species and Pielow's evenness index has presented significant differences after clipping treatment in the localities of Mallorca (Aucanda and Es Barcares). A large number of factors and

phenomena can alter the structure of the epifauna community and, therefore, its resilience. Due to this complexity, it is difficult to know the role of herbivory pressure in the regulation of epifauna community. Even so, replications made at different geographical scales have shown that herbivory pressure is present in the variations that this type of ecosystem can suffer. It should be pointed out that the changes in the epifauna community due to the herbivory pressure differs depending on the location. For this reason, it is important to perform more studies in different locations to be sure that representative patterns are obtained. Altogether, this work emphasize the importance of taking into account the herbivory pressure as factor to investigate in subsequent studies that are carried in *C. nodosa* meadows.

6. Ethics and morality of the project

In the daily activity, carried put in the laboratory of Instituto Mediterráneo de Estudios Avanzados (IMEDEA) located in Esporlas (Mallorca), the safety regulations have been complied with. In addition, personnel have been informed of the management of both chemical and daily waste. In this laboratory, waste reduction has been carried out using recycled materials, when have been possible.

In the fieldwork, precautions have been taken to avoid any damage during the dives. In addition, maintenance has been carried out on the material used during, before and after the dives.

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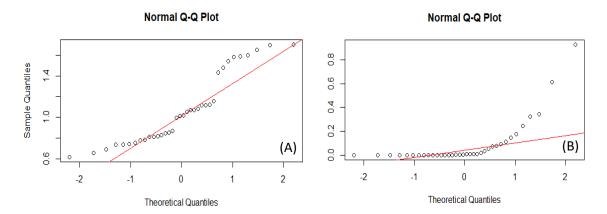
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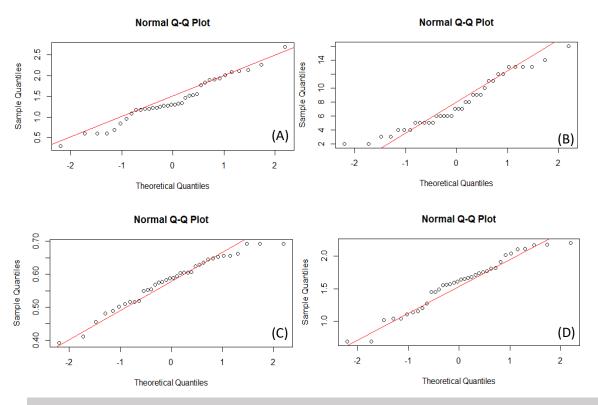
9. Appendix



S1. Normal Q-Q plot from biomass of C. nodosa (A) and biomass of epiphytes (B)

S2. Wilcoxon sum-rank test (U-Mann Whitney) as a post-hoc analysis according to different region fixed factor, Gran Canaria and Mallorca; and the response biomass of *C. nodosa* (gDW/100cm²) and biomass of epiphytes (gDW/100cm²). Significant p-values are highlighted in bold.

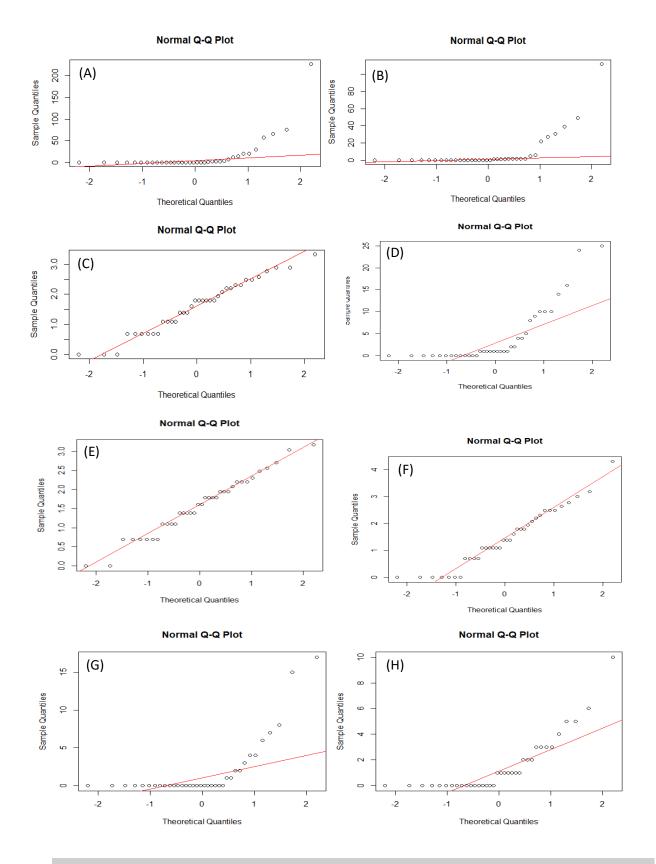
Source of variation			W	p-value
Biomass of Cymodocea				
nodosa (gDW/100cm²)				
Ca	astillo			
		Control*Cut80%	20	0.01587
Ar	rinaga			
		Control*Cut80%	20	0.01587
Aud	canda			
		Control*Cut80%	18	0.06349
Es Bar	rcares			
		Control*Cut80%	16	0.1905
Biomass of epiphytes				
(gDW/100cm ²)				
Ca	astillo			
		Control*Cut80%	10	NA
Ar	rinaga			
		Control*Cut80%	12.5	0.5605
Aud	canda			
		Control*Cut80%	19	0.03175
Es Bar	rcares			
		Control*Cut80%	11	0.9048



S3. Normal Q-Q plot from abundance of individuals (A), taxa richness (B), Pielow's evenness index (C) and Shannon Wiener index (D).

S4. Tukey HSD test as a post-hoc analysis according to different region fixed factor, Gran Canaria and Mallorca; and the response of abundance of individuals (ind/100cm²), taxa richness (species/100cm²), Pielow's evenness index (J') and Shannon Wiener index (H').

Source of variation			diff	p-adj
Abundance (ind/100cm²)				
	Castillo	Control*Cut80%	0.22064702	0.0011110
	Arinaga	Control*Cut80%	-0.22964702	0.9911119
	, ii ii iaga	Control*Cut80%	-0.01035836	1.0000000
	Aucanada			
		Control*Cut80%	0.37449860	0.8833004
	Es Barcares	Control*Cut900/	0.61450207	0.4001636
Taxa richness		Control*Cut80%	-0.61450397	0.4001636
(species/100cm²)				
	Castillo			
		Control*Cut80%	0.0375000	1.0000000
	Arinaga	Control*Cut80%	0.4635000	0.0000083
	Aucanada	Control Cut80%	-0.4625000	0.9999982
	Addanada	Control*Cut80%	1.6125000	0.9930633
	Es Barcares			
		Control*Cut80%	-2.0375000	0.9734540
Pielow's eveness (J)	Ca atilla			
	Castillo	Control*Cut80%	0.030920401	0.9929853
	Arinaga	Control Catoo70	0.030320401	0.5525055
		Control*Cut80%	-0.01524053	0.9999263
	Aucanada			
	F. B	Control*Cut80%	-0.07640933	0.5475214
	Es Barcares	Control*Cut80%	0.09208919	0.3194721
Shannon Wiener (H')		Control Cuto0/0	0.03200313	0.3134721
	Castillo			
		Control*Cut80%	-0.00247750	1.0000000
	Arinaga	0 1 140 1000	0.40507753	0.00000.55
	Aucanada	Control*Cut80%	-0.19597750	0.9939946
	Aucandua	Control*Cut80%	-0.03987250	0.9999999
	Es Barcares			
		Control*Cut80%	-0.15858250	0.9983836



S5. Normal Q-Q plot from abundance of Nematoda (A), Polychaeta (B), Prosobranchia (C), Caprellidea (D), Gammaridea (E), Copepoda (F), Cumacea (G) and Tanaidacea (H).

S6. Wilcoxon sum-rank test (U-Mann Whitney) as a post-hoc analysis according to different region fixed factor, Gran Canaria and Mallorca; and the response of abundance of Nematoda, Caprellidea, Polychaeta, Cumacea and Tanaidacea taxa.

Source of variation			W	p-value
Nematoda				
	Gran Canaria			
		Control*Cut80%	45	0.3741
	Mallorca			
		Control*Cut80%	38.5	0.894
Caprellidea				
	Gran Canaria			
		Control*Cut80%	46.5	0.5715
	Mallorca			
		Control*Cut80%	56.5	0.1689
Polychaeta				
	Gran Canaria			
		Control*Cut80%	38.5	0.8667
	Mallorca	0 1 1 2 1 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	27	0.7067
		Control*Cut80%	37	0.7867
Cumacea	Cara Cara ta			
	Gran Canaria	C	40.5	210
	Mallana	Control*Cut80%	40.5	NA
	Mallorca	Comtral*C+000/	40	1
Tanaidassa		Control*Cut80%	40	1
Tanaidacea	Cran Canaria			
	Gran Canaria	Control*Cut80%	37.5	0.7911
	Mallorca	Control*Cut80%	37.5	0.7911
	ivialiofCd	Control*Cut80%	34.5	0.6218
		Control Cutou/6	٠	0.0210

S7. Tukey HSD test as a post-hoc analysis according to different region fixed factor, Gran Canaria and Mallorca; and the response of abundance of Prosobranchia and Copepoda taxa.

Source of variation			diff	p-adj
Prosobranchia				
	Gran Canaria			
		Control*Cut80%	-0.8760326	0.1290929
	Mallorca			
		Control*Cut80%	-0.2898703	0.8751600
Copepoda				
	Gran Canaria			
		Control*Cut80%	-0.3439163	0.7674031
	Mallorca			
		Control*Cut80%	-0.1797644	0.9567442