



Fast climatic changes place an endemic Canary Island macroalga at extinction risk

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Abstract

Global warming and ultraviolet radiation (UVR) affect marine organisms worldwide. However, we still lack knowledge about the consequences of these environmental changes to range-restricted macroalgae species, such as *Gelidium canariense*, a habitat-forming endemic of the Canary Islands. The aim of this study is to provide useful data for analyzing the conservation status of this endemic species. To do this, (1) we compared distributional data from 1987 with the data collected in 2008 and 2019 along the northern coast of Tenerife; (2) we assessed the variation in macroalgae assemblages dominated by *G. canariense* over the last decade on this same coast; and (3) we explored the correlation of these results with the actual trends of sea surface temperature (SST), air temperature at 2 m above the sea surface (T2m), UVR, and downward solar radiation (DSR). The results showed significant changes in the distribution of *G. canariense* in the last 30 years at Tenerife. More than 90% of its populations have declined. The endemic macroalgae have been replaced by the globally distributed species *Pterocladia capillacea* at nearly all the sites studied. The cover of *G. canariense* and *Gelidium arbuscula* decreased significantly with increasing SST and T2m. Additionally, *G. canariense* is negatively related to the rise in UVR. These results highlight the vulnerability of both species of *Gelidium* to global warming and UVR, but especially of *G. canariense*. The drastic decline of this species reveals the need to upgrade its conservation status and implement protection measures to reduce the damage that predicted global change scenario may have to their relict populations.

Keywords Canary Islands · Decline · *Gelidium* · Range-restricted · SST · Ultraviolet radiation

Introduction

Global warming and ultraviolet radiation (UVR) can affect marine and terrestrial ecosystems worldwide (Myers et al. 2000; Parmesan and Yohe 2003; Schiel et al. 2004; Bischof et al. 2006; Mieszkowska et al. 2006; Hawkins et al. 2009;

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Ballaré et al. 2010; Burrows et al. 2011; Morueta-Holme et al. 2015). To countless organisms, the rise in ocean and air temperatures can result in distributional shifts, abundance or density changes, phenological modifications (i.e. migration, flowering, and spawning), morphological changes (i.e. body size), and even genetic frequency shifts (Root et al. 2003; Parmesan and Yohe 2003; Both et al. 2009; La Sorte and Jetz 2010; Casado-Amezúa et al. 2019). Additionally, the effects of UVR may reduce the uptake of atmospheric carbon dioxide in photosynthetic organisms, generate DNA damage, and affect species diversity, ecosystem stability, trophic interactions and global biogeochemical cycles (Häder et al. 2011; Schimitz et al. 2018). Likewise, the negative effects of UVR may be augmented by other environmental changes, including global climate change processes (Bais et al. 2015; Andradý et al. 2017; Barnes et al. 2019).

Species vulnerability to global warming and changes in UVR may depend on both ecological and geographical characteristics (Broennimann et al. 2006) and the ability of organisms to adapt to new conditions, successfully disperse and migrate to existing suitable habitats. Among seaweeds, temperature delimits three different types of essential boundaries: lethal boundaries (i.e., the ability of species to survive unfavorable seasons), growth boundaries (i.e., the temperature range in which species can grow), and reproductive boundaries (i.e., the temperature range suitable for reproduction) (Bartsch et al. 2012; Eggert 2012). These boundaries are generally species-specific; however, characteristics of the ecosystem that the species inhabits play a fundamental role in determining species adaptability. For example, intertidal macroalgae already live in an extremely dynamic environment and are adapted to the stress of emersion, while changes in irradiance and desiccation are more stressful for subtidal species (Bischof et al. 1998; Wiencke et al. 2006). Additionally, the successful dispersal of macroalgae by propagules, such as spores or zygotes, is lower than that in terrestrial plants due to differences in the surrounding environment and motility and viability of their propagules (Santelices 1990; Norton 1992; Gaylord et al. 2002, 2006; Incagnone et al. 2015). Ultimately, the potential of species to adapt to new conditions, together with their dispersal capability, determines their ability to migrate to existing habitats, if available.

Furthermore, the ability of species to withstand environmental changes (e.g., increasing temperature and UVR) will differ between wide-ranging species and range-restricted (endemic) species. The latter live under only limited environmental conditions and would thus struggle more to adapt to new conditions and may disappear when lethal boundaries are exceeded. Marine macroalgal endemism is less common than terrestrial plant endemism. Some endemic marine macroalgae play major

roles in certain ecosystems, and during the last decade, studies have focused mainly on the risks that global warming may pose to these species. The direct consequences of rising temperatures for endemic species are, on the one hand, population reductions and even local extinctions (Thibaut et al. 2005; Smale and Wernberg 2013; Buonomo et al. 2018). On the other hand, detrimental effects of warming on early life stages have been reported for several species (Andrews et al. 2014; Wernberg et al. 2016; Capdevila et al. 2019). Additionally, indirect consequences arise when endemic macroalgae are the dominant species in an ecosystem; hence, the loss of habitat-forming species would impact other species of the benthic community (Smale and Wernberg, 2013; Capdevila et al. 2016). In contrast, deleterious effects of UVR on endemic macroalgae have rarely been studied. Only Arctic and Antarctic endemic macroalgae species have been thoroughly studied in relation to increasing UVR (Karsten et al. 2003; Roleda et al. 2007a, b; Houvinen and Gómez 2013; Häder 2018), while the remaining studies have focused on widespread or important commercial macroalgae species (Gomez and Figueroa 1998; Bischof et al. 2006; Navarro et al. 2019; Quintano et al. 2017). Nevertheless, all of the previous studies agree that changes in UVR can shape seaweed communities (e.g., by conditioning the vertical zonation) of macroalgae and that most certainly, early developmental stages are the most vulnerable to the increase in UVR.

Nonetheless, a distinction should be made between continental and island endemic seaweeds. In general, islands are considered gathering places for algae propagules from different biogeographical areas (Sansón et al. 2001), while they are treated as areas of isolation for terrestrial species (Whittaker et al. 2007). However, algae endemic to islands are more susceptible to dying off and eventually result in global extinction because the migration is limited to the island's boundaries. The total number of species on islands is generally lower than that on continents; nevertheless, the high level of endemism in islands at all taxonomic levels highlights their extensive contribution to global biodiversity (Malcolm and Markham 2000; Wilmé et al. 2006; Whittaker and Fernández-Palacios 2007; Kier et al. 2009; Veron et al. 2019). Despite this fact and the inherent vulnerability of islands' endemic macroalgae, studies of the direct or indirect effects of global warming and UVR on these species are scarce. Some studies have focused on short- and long-distance dispersal pathways of an endemic fucal species of New Zealand (Buchanan and Zucarello 2012); others have unraveled the extensive cryptic species diversity and the fine-scale endemism of *Portieria* species in the Philippine archipelago (Payo et al. 2012). However, we still lack knowledge about the consequences of climate

change on this kind of vulnerable endemic seaweeds and the indirect implications at the community level after their disappearance.

The Canary Islands (28° N, 16° W) are an oceanic archipelago situated in the subtropical gyre of the northeastern Atlantic Ocean. The upwelling of cold waters off the coast of northwestern African and the warmer waters from the open ocean influence these islands (Barton et al. 1998; Van den Bogaard 2013). These attributes create a sea surface temperature gradient of almost 2 °C from the eastern to the western islands of the archipelago (Barton et al. 1998). Ultimately, this generates fringe habitats that hamper species from migrating between islands and makes any dispersion to the nearest continent more difficult. Among the marine flora of the archipelago are approximately 690 algal species (Gallardo et al. 2016), of which approximately 31% are restricted to the northeastern Atlantic warm-temperate region and only 3% (i.e., 24 species) are exclusive endemic algae from the Canary Islands (Sansón et al. 2001).

Gelidium canariense is the only endemic and habitat-forming macroalga of the archipelago. This rhodophyte is a dominant species in the upper subtidal zone of the northern rocky shores of four central and western islands. According to historical observational data, populations of *G. canariense* have suffered an extensive decline in recent decades (Martínez et al. 2015); however, no previous studies have supported this observation with accurate data or characterized the relationship of this species with environmental changes. Hence, the goal of this research is to provide useful data for analyzing the status of this species, which is relevant to global biodiversity. In this study, (1) we evaluated changes in the distribution of the endemic macroalga *Gelidium canariense* during the last 30 years at the northern shore of Tenerife, (2) we determined the variation in macroalgae assemblages dominated by *G. canariense* over the last decade on this same coast, and (3) we explored whether these changes correlate with the actual trends of some climatic variables in a scenario of global environmental change.

Material and methods

Study site

The Canary Islands are a volcanic archipelago constituted by 7 major islands, several smaller islands and islets and several submarine seamounts. The environmental conditions, together with the geographical position of these islands, contribute to the presence of a unique marine flora. Species of *Gongolaria*, *Ericaria*, *Cystoseira*, Dictyotales, Gelidiales, geniculate coralline algae (GCA), and crustose coralline algae (CCA) and a mix of turf-forming species compose

the main macroalgae assemblages on the rocky shores of the archipelago (Sangil et al. 2012, 2018). The only habitat-forming endemic species in the archipelago is *Gelidium canariense* (Alfonso et al. 2019). This perennial red alga inhabits only the upper limit of the subtidal zone along the northern shores of four islands: La Palma, La Gomera, Tenerife, and Gran Canaria. Due to its restricted distribution and observed reductions in its populations, *G. canariense* has been recently classified as a vulnerable species in the *List of Wild Species under Special Protection Regime and Spanish Catalog of Threatened Species* (Orden TEC/596/2019, BOE 2019). The study area is located on the northern rocky shore of Tenerife, where the largest populations of *G. canariense* are found. This species occurs mainly with the two other habitat-forming Gelidiales: *Pterocladia capillacea* and *Gelidium arbuscula*, which usually grow in greater abundance just above *G. canariense*.

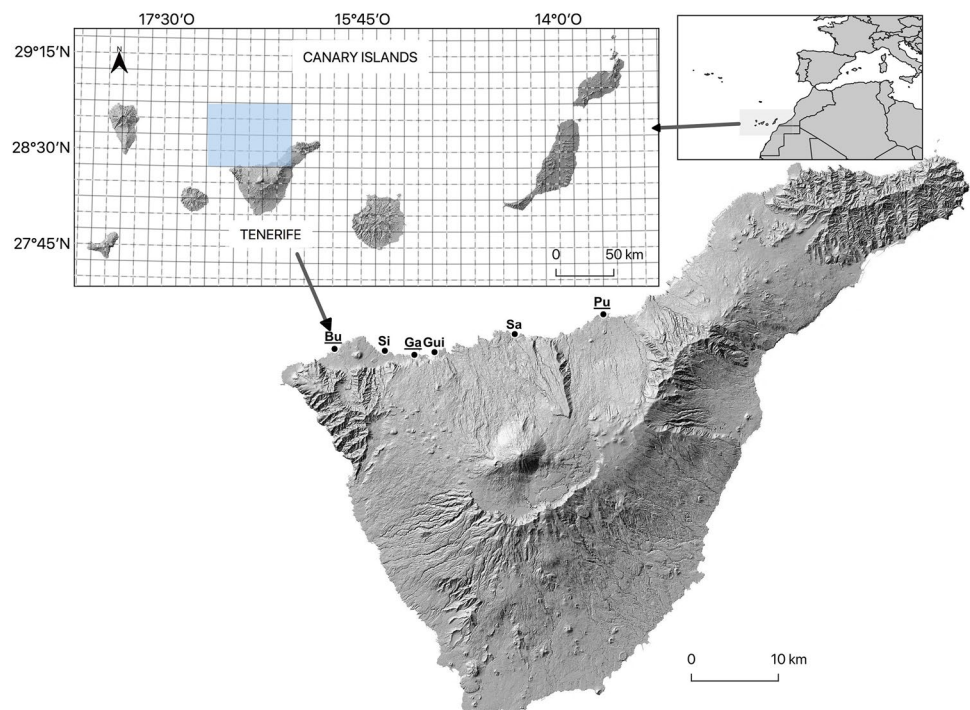
Historical changes in distribution

The first data on the spatial distribution of assemblages dominated by *Gelidium canariense* were documented by Wildpret et al. (1987) as part of a broader characterization of the marine macroalgae communities in the Canary Islands. Subsequently, Rodríguez et al. (2008) provided distributional data, adding coverage data of the assemblages dominated by *G. canariense*. For the current study, we randomly selected 3 sites in the former distribution of *G. canariense* in Tenerife (Buenavista, Garachico, and Puerto de la Cruz) (Fig. 1). The presence and absence of *G. canariense* was recorded along a minimum of 2 km per site. The total distribution obtained was divided into sections of 100 m in length at each site and compared among years. The data were collected using MapitGIS (spatial asset collection, GPS surveys, and data management for android) and exported into the free and open source Geographic Information System QGIS. The results are expressed as the average extension in meters per section \pm SE.

Macroalgae assemblage variation

To evaluate the variation over the last decade in assemblages dominated by *G. canariense*, the percentage of cover of all macroalgae species was estimated in 2017 and 2019 at six of the sites previously studied by Rodríguez et al. (2008) in Tenerife (Buenavista, Los Silos, Garachico, El Guincho, San Juan, and Puerto de la Cruz) (Fig. 1). To achieve comparable data sets between years, the same methodology used in Rodríguez et al. (2008) was followed. However, to obtain a better representation of these assemblages and to improve the long-term monitoring program, in 2017 and 2019, we surveyed a total of 6 transects of 10 m and 3 photocuadrats of 50 × 50 cm per transect along the *G. canariense* belt per

Fig. 1 Study areas on the north shore of Tenerife (Canary Islands, NE Atlantic). Underlined sites indicate areas where historical distribution was assessed (Bu = Buenavista, Ga = Garachico and Pu = Puerto de la Cruz). Variation in macroalgae assemblages was studied at all sites (Bu, Si = Los Silos, Ga, Gui = El Guincho, Sa = San Juan and Pu). Climatic variables data were obtained for the blue square grid off the north coast of Tenerife



site, instead of 3 transects as performed in Rodríguez et al. (2008). The percentage of cover of the different macroalgae species in each photoquadrat was obtained using the open source image processing program ImageJ. To minimize differences between years and prevent misidentifications, some species were aggregated into genera or, corresponding to morphological classifications, turf-forming species, crustose coralline algae (CCA), geniculate coralline algae (GCA), and cyanobacteria, according to Littler and Littler (1980). We defined turf-forming species as sparse to thick mats of diminutive and juvenile algae that are < 2 cm tall (Littler and Littler 2011).

Climatic variables

Monthly averages for the period 1979–2018 of sea surface temperature (SST), air temperature at 2 m above the surface of the seawater (T2m), downward UV radiation at the surface (UVR), the amount of ultraviolet radiation reaching the surface with a wavelength of 0.20–0.44 nm, and downward solar radiation (DSR) the amount of solar radiation (i.e., shortwave radiation) that reached the surface, corresponding to the northern shore of Tenerife Island were obtained from the Copernicus Climate Change Service (European Commission) with a horizontal grid resolution of $0.25^\circ \times 0.25^\circ$ (Fig. 1). These data correspond to ERA5, the last climate reanalysis dataset produced by the European Centre for Medium Range

Weather Forecasts (ECMWF, <https://www.ecmwf.int>). The reanalysis data combine observations from across the world with model data. This combination permits us to estimate biases between observations and produce data where in situ coverage is low. One of the novelties in the ERA5 dataset is the estimation of uncertainty using the 10-member data assimilation (EDA) system. Although the reanalysis data from ERA-5 are designed for global studies, the only meteorological station located at the study site is situated 2000 m above sea level (Izaña Atmospheric Research Centre–AEMET); hence, we considered the data from ERA-5 very useful in the absence of any other available source to obtain long-term climatic variables at sea level.

Data analysis

Historical changes in the distribution of *G. canariense* were analyzed by performing a permutational analysis of variance (ANOVA) (Anderson et al. 2008) with a two-factor design that included “year” (random, 3 levels) and “site” (random, 3 levels). Likewise, a permutational ANOVA was used to test the variability in the macroalgae in assemblages dominated by *G. canariense* among years and sites. The design included two factors: “year” (random, 3 levels) and “site” (random, 6 levels). Both analyses were performed on a Bray–Curtis similarity matrix calculated on square-root transformed data. Because the design of the assemblage

Table 1 Permutational analysis of variance (ANOVA) of the distribution of *Gelidium canariense* in the north shores of Tenerife, with factors 'year' (random, 3 levels) and 'site' (random, 3 levels)

Source	df	SS	MS	Pseudo-F	P(perm)
year	2	92427	46213	1.761	0.29
site	2	50249	25125	0.942	0.434
year × site	4	106000	26652	18.23	0.000
Residual	243	355000	1462		
Total	251	620000			

variation analysis was unbalanced, to obtain more power, the total number of samples from 2017 and 2019 at each site ($n = 18$) was randomly reduced to half ($n = 9$), and the same statistical analysis was performed again. If different results were obtained, this would point out that differences in the number of replicates between years affected the analysis. This was done three times with three different sets of random samples. To identify the macroalgae species that contribute most to the variability among "years" and "sites," a SIMPER analysis was performed. All the statistical analyses were performed using PRIMER 7 and PERMANOVA+ (Anderson et al. 2008) software and statistical package.

The temporal trend of each climatic variable was evaluated by calculating the linear regression of the anomalies of each variable *versus* the time for the total period per year. The climatic variables with significant regressions were related to the cover values of the macroalgae species that contributed most to shifts in the last decade, according to the SIMPER analysis. The analysis was performed with the IBM SPSS 26 statistical package.

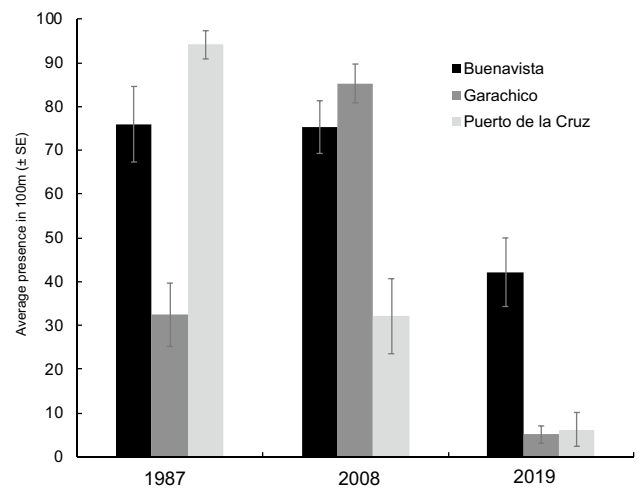
Results

Historical changes in distribution

The distribution of *Gelidium canariense* has changed significantly over the last 3 decades (Table 1) at the north rocky shores of Tenerife. Puerto de la Cruz, the populations of which occupied the entire coastline in 1987,

Table 2 Permutational analysis of variance (ANOVA) of the percentage of cover of macroalgae in *Gelidium canariense* assemblages, with factors 'year' (random, 3 levels) and 'site' (random, 6 levels)

Permutational ANOVA					
Source	df	SS	MS	Pseudo-F	P(perm)
year	2	32924	16462	5.110	0.002
site	5	46389	9277.8	4.089	0.000
year × site	10	32211	3221.1	4.112	0.000
Residual	216	169100	783.19		
Total	233	353000			

**Fig. 2** Average presence (meters \pm SE) of *Gelidium canariense* along a transect of 100 m examined at each site and year. Significant interaction of site and year was obtained in the permutational ANOVA analysis

suffered the greatest decline in *G. canariense* extension, with average reductions from $94.07 \text{ m} \pm 3.21 \text{ m}$ (\pm SE) to $6.32 \text{ m} \pm 3.85$ over 100 m of coast, which corresponded to a 93% reduction in the previous distribution. Similarly, at Garachico, the decrease in the population ranged from $32.47 \text{ m} \pm 7.22$ to 5.13 ± 1.97 , for a loss of 84% of the original population. Finally, Buenavista currently harbors the most extensive distribution. In this location, the decrease in the population ranged from $75.95 \text{ m} \pm 8.62$ to $42.17 \text{ m} \pm 7.83$, which corresponds to a reduction of 45% of the former populations. The results also show that the greatest changes in populations along the island have taken place over the last decade (Fig. 2) and that the populations that remain are conserved as fragmented populations.

Macroalgae assemblage variation

The analysis of macroalgae species in the assemblages dominated by *G. canariense* shows significant variation in

Table 3 SIMPER analysis of species that contribute up to the 70% of the dissimilarity among levels of factors ‘year’ and ‘site’. CCA, crustose coralline algae; GCA, geniculate coralline algae; G. arb. *Gelidium arbusculum*; G. can, *Gelidium canariense*; P. cap, *Pterocladia*

capillacea; Turf, turf-forming species. * Denotes species that contribute more than 60% of the dissimilarities among groups of ‘year’ and ‘site’. Bu, Buenavista; Si, Los Silos; Ga, Garachico; Gui, El Guincho; Sr, San Juan; Pu, Puerto de la Cruz

SIMPER		Year		Site							
Species	2008 vs. 2017		Species	2008 vs. 2019		Species	2017 vs. 2019				
	Cont%	Cum%		Cont%	Cum%		Cont %	Cum%			
P. cap*	21.58	21.58	P. cap*	18.57	18.57	P. cap*	16.53	16.53			
G. can*	19.28	40.87	G. can*	17.63	36.20	G. can*	15.34	31.87			
CCA*	17.86	58.73	G. arb*	14.58	53.01	CCA*	14.64	46.51			
G. arb*	17.32	76.05	CCA*	16.80	67.59	Turf*	13.40	59.91			
			Turf*	7.69	75.28	G. arb*	12.45	72.36			
	Av. dis. = 56.60			Av. dis. = 61.64			Av. Dis. = 40.44				
Site		Bu vs. Si		Bu vs. Ga		Si vs. Ga		Bu vs. Gui			
	Cont%	Cum%		Cont%	Cum%		Cont%	Cum%		Cont%	Cum%
P. cap*	22.85	22.85	P. cap*	30.83	30.83	P. cap*	26.71	26.71	G. can*	28.81	28.81
G. can*	11.63	41.96	CCA*	18.87	49.70	G. can*	22.35	49.05	CCA*	18.07	46.88
CCA*	19.11	55.80	G. can*	16.85	66.55	CCA*	9.77	58.82	G. arb*	16.97	63.85
G. arb*	13.83	67.43	G. arb*	11.45	78.01	G. arb*	7.64	66.46	P. cap*	15.78	79.62
Codium	10.63	78.06				Codium	7.58	74.04			
	Av. dis. = 47.70			Av. dis. = 62.12			Av. dis. = 61.38			Av. dis. = 46.82	
Si vs. Gui		Ga vs. Gui		Bu vs. Sr		Si vs. Sr					
	Cont%	Cum%		Cont%	Cum%		Cont%	Cum%		Cont%	Cum%
CCA*	19.31	19.31	P. cap*	31.52	31.52	G. can*	22.19	22.19	CCA*	14.88	14.88
P. cap*	18.74	38.05	G. can*	24.67	56.18	P. cap*	15.01	37.20	P. cap*	14.18	29.06
Codium	13.59	51.64	CCA*	12.87	69.06	G. arb*	14.40	51.60	Turf*	12.33	41.39
G. arb*	11.56	63.20	G. arb*	8.32	77.37	Turf*	13.95	65.60	GCA	10.94	52.34
Turf*	10.99	74.19				CCA*	12.12	77.67	G. arb*	10.54	62.87
	Av. Dis. = 39.61			Av. Dis. = 70.62			Av. Dis. = 48.63		Codium	10.23	73.10
										Av. Dis. = 47.22	
Ga vs. Sr		Gui vs. Sr		Bu vs. Pu		Si vs. Pu					
	Cont%	Cum%		Cont%	Cum%		Cont%	Cum%		Cont%	Cum%
P. cap*	24.44	24.44	P. cap*	17.46	17.46	G. can*	24.94	24.94	CCA*	17.85	17.85
G. can*	19.27	43.71	CCA*	16.92	34.32	Turf*	19.97	44.90	Turf*	15.76	33.61
CCA*	13.92	57.63	Turf*	14.06	48.44	G. arb*	16.03	60.93	G. can*	13.84	47.46
Turf*	8.49	66.12	GCA	13.35	61.79	P. cap*	15.09	76.03	P. cap*	13.69	61.15
G. arb*	8.41	74.53	G. arb*	11.38	73.17				Codium	10.72	71.82
	Av. dis. = 65.96			Av. dis. = 44.50			Av. dis. = 42.94			Av. dis. = 46.95	
Ga vs. Pu		Gui vs. Pu		Sr vs. Pu							
	Cont%	Cum%		Cont%	Cum%		Cont%	Cum%			
P. cap*	26.71	26.71	Turf*	20.43	20.43	Turf*	18.04	18.04			
G. can*	18.05	44.75	P. cap*	18.94	39.38	P. cap*	16.25	34.29			
CCA*	16.43	61.19	G. can*	17.65	57.02	G. can*	15.65	49.94			
Turf*	11.86	73.05	CCA*	16.87	73.89	GCA	12.79	62.74			
	Av. dis. = 67.94			Av. dis. = 42.06		CCA*	10.98	73.71			
							Av. dis. = 46.65				

the interaction term of the random factors (“year × site”) (Table 2). The percentage of macroalgae cover has significantly changed in the last decade. The SIMPER analysis revealed that *G. canariense*, *G. arbuscula*, *P. capillacea*,

coralline crustose algae (CCA), and turf-forming species explain more than 60% of the total spatial and temporal variation in macroalgae species cover (Table 3). In 2008, *G. canariense* was the dominant species in Los Silos,

Garachico, and San Juan, with a mean percentage cover (\pm SE) of 57.22 ± 9.63 , 46.22 ± 8.12 , and 36.11 ± 14.28 , respectively. Ten years later, this endemic species had almost disappeared in Los Silos ($1.68 \pm 0.85\%$) and San Juan ($9.97 \pm 3.69\%$), whereas in Garachico, 60% of the species cover remained. The small population of *G. canariense* documented in El Guincho in 2008 disappeared completely in 2017. *G. arbuscula* was the dominant species in Buenavista ($60.55 \pm 9.34\%$) and El Guincho (17.77 ± 2.42), although it has drastically declined in the last decade at both sites (Buenavista: $17.14 \pm 4.63\%$ in 2017 and $3.57 \pm 1.71\%$ in 2019; El Guincho: 2.36 ± 1.72 in 2019). In 2008, *Pterocliadiella capillacea* was the dominant species in assemblages of Puerto de la Cruz only, but since 2017, it has become the dominant species at all studied sites of the island, with abundances above 50%, except for Garachico, where abundances have remained low. CCA only contributed to the macroalgae assemblages in San Juan and Puerto de la Cruz in 2008, while in the last decade, they have become an important contributor in Garachico ($21.46 \pm 3.67\%$), El Guincho ($18.45 \pm 4.11\%$), Los Silos ($3.13 \pm 0.93\%$), and Buenavista ($3.59 \pm 2.51\%$). Similarly, turf-forming species were not present 10 years ago; however, they are currently inhabiting all sites of the island, with the highest abundances in San Juan ($18.04 \pm 3.52\%$) and Puerto de la Cruz ($11.78 \pm 3.77\%$) and the lowest in Los Silos ($2.25 \pm 0.82\%$) and Buenavista ($0.023 \pm 0.02\%$) (Table 4; Figs. 3 and 4).

Importance of climatic variables to macroalgae assemblage variation

The SST, T2m, and UVR have increased significantly over the last 40 years off the northern coast of Tenerife. SST and T2m have risen 0.2 °C per decade, whereas UVR has increased 0.1 W/m² per decade. DSR has shown no significant differences over the last 4 decades (Fig. 5). For the *Gelidium* species, the relationships of the mean percentage of cover to the climatic variables show significant linear regressions. The cover of *G. canariense* decreased significantly in the last decade in parallel with the increase in SST, T2m, and UVR (Fig. 6a), whereas the reduction in cover of *G. arbuscula* is related to rises in SST and T2m (Fig. 6b). Regarding *P. capillacea*, no significant relationship with the climatic variables was observed (Fig. 6c), despite having an evident increase of nearly 30% in cover in the last decade. The CCA and turf-forming species did not show significant changes with the trends of the climatic variables (Figs. 6d and e).

Discussion

For the first time, the reduction in the population of a macroalga endemic to oceanic islands over the last 30 years is demonstrated. This study revealed a negative correlation between the cover of the endemic and habitat-forming species *G. canariense* from the Canary Islands, the global warming and the increase in ultraviolet radiation over the last decade in Tenerife Island. Furthermore, the results show that indirect consequences of its decrease have resulted in changes at the assemblage level, creating available niche space for *P. capillacea* to proliferate and causing local extinctions at some sites within its range of distribution in this island.

Temperature is a major factor controlling the rate of photosynthesis and growth in all plants and algae (Davison 1991; Clark et al. 2013), and in *G. canariense* and *G. arbuscula*, the results suggest that rising air and ocean temperatures are important factors determining their presence and cover in the archipelago. These species inhabit the upper limit of the subtidal zone, so they are exposed to air during low tides and are submerged under the surface of the water for the rest of the time (Polifrone et al. 2012; Alfonso et al. 2017, 2019). Consequently, their decrease in cover with the rise in SST and T2m suggest that the species are sensitive to warming and that are vulnerable to changes in seawater but also air temperature during emersion periods. In fact, previous experimental studies have demonstrated the relation of the vertical zonation of these species to their low tolerance to emersion (Domínguez-Álvarez et al. 2011).

In addition, increasing UVR was negatively correlated with a significant decrease only in *G. canariense* cover. This endemic species grows only in the northern rocky shores of La Palma, La Gomera, Tenerife, and Gran Canaria islands, where there is the greatest accumulation of clouds carried by trade winds (Martin et al. 2011). Mercado et al. (2001) demonstrated that the photosynthetic rate of this species might be limited by incident light. This could suggest an adaptation to a narrow and species-specific level of UVR and would explain why this endemic species grows on only the northern rocky shores of the islands, while the other two habitat-forming Gelidiales species can grow along the coasts throughout the archipelago. It has been determined that in many macroalgae, the most UV-susceptible life stages are their propagules (Roleda et al. 2007a, b; Wiencke and Amsler 2012; Navarro et al. 2016, 2019). Hence, the finding that UVR was only negatively correlated with the endemic species cover, may suggest not only an adaptation of the adults of this species to its habitat but also the vulnerability of its early life stages to changes in this parameter.

Table 4 Mean percentage cover of macroalgae (% ± SE) registered in each year and site at Tenerife

	Species	Buenavista	Los Silos	Garachico	El Guincho	San Juan	Puerto Cruz
2008	Rhodophyta						
	<i>Corallina</i> sp.	-	-	-	13.33 ± 0.00	-	-
	Crustose coralline algae	2.22 ± 1.47	-	-	-	20.00 ± 10.71	13.89 ± 3.89
	<i>Gelidium arbuscula</i>	60.55 ± 9.34	6.89 ± 1.56	4.11 ± 0.55	17.77 ± 2.42	22.22 ± 2.22	18.78 ± 4.47
	<i>Gelidium canariense</i>	26.66 ± 6.94	57.22 ± 9.63	46.22 ± 8.12	2.22 ± 0.55	36.11 ± 14.28	21.66 ± 3.46
	<i>Laurencia</i> sp.	7.22 ± 7.22	-	-	-	-	-
	<i>Pterocladia capillacea</i>	-	15.00 ± 0.96	5.55 ± 3.88	14.44 ± 4.01	18.89 ± 1.11	35.55 ± 5.88
	Phaeophyceae						
	<i>Halopteris scoparia</i>	2.77 ± 2.77	-	-	-	-	-
	<i>Sargassum</i> sp.	-	-	-	-	-	0.55 ± 0.55
	<i>Gongolaria abies-marina</i>	-	-	0.33 ± 0.33	-	-	-
	Chlorophyta						
	<i>Codium</i> sp.	-	1.11 ± 1.11	-	1.11 ± 0.55	-	-
	<i>Ulva</i> sp.	-	-	-	-	0.33 ± 0.33	7.22 ± 4.33
Cyanobacteria	-	-	-	2.22 ± 2.22			
2017	Rhodophyta						
	<i>Corallina</i> sp.	0.36 ± 0.35	-	-	-	3.51 ± 1.66	2.83 ± 2.38
	Crustose coralline algae	4.78 ± 3.14	20.70 ± 3.27	29.42 ± 5.28	6.12 ± 1.45	9.87 ± 2.98	1.44 ± 0.58
	<i>Gelidium arbuscula</i>	17.14 ± 4.63	2.87 ± 0.70	6.22 ± 2.19	-	5.71 ± 2.02	3.01 ± 1.36
	<i>Gelidium canariense</i>	22.90 ± 6.03	2.15 ± 1.53	42.42 ± 6.62	-	9.97 ± 3.69	12.38 ± 4.46
	<i>Grateloupia imbricata</i>	-	-	0.02 ± 0.02	-	-	-
	<i>Laurencia</i> sp.	-	0.85 ± 0.27	-	0.15 ± 0.15	2.47 ± 1.03	1.73 ± 1.34
	<i>Pterocladia capillacea</i>	51.36 ± 4.78	48.22 ± 5.12	3.72 ± 2.03	78.86 ± 4.77	54.60 ± 6.52	52.65 ± 6.80
	Phaeophyceae						
	<i>Sargassum</i> sp.	-	-	0.39 ± 0.21	-	-	-
	<i>Padina pavonica</i>	-	-	0.004 ± 0.004	-	-	-
	Chlorophyta						
	<i>Anadyomene saldanhae</i>	-	0.02 ± 0.2	-	-	-	-
	<i>Chaetomorpha</i> sp.	-	0.03 ± 0.02	-	-	-	-
	<i>Cladophora</i> sp.	-	-	-	-	0.007 ± 0.007	-
	<i>Codium</i> sp.	0.02 ± 0.02	5.20 ± 1.37	0.34 ± 0.21	-	-	-
	<i>Ulva</i> sp.	-	0.20 ± 0.09	0.08 ± 0.03	-	-	-
	Cyanobacteria	-	0.39 ± 0.35	0.60 ± 0.28	0.37 ± 0.22	1.03 ± 0.56	0.04 ± 0.04
	Turf	0.61 ± 0.44	3.32 ± 1.28	1.04 ± 0.33	3.50 ± 1.73	4.95 ± 0.79	18.33 ± 5.56
	2019	Rhodophyta					
<i>Asparagopsis taxiformis</i>		-	-	-	-	0.11 ± 0.11	-
<i>Corallina</i> sp.		-	-	3.22 ± 3.22	-	14.89 ± 4.73	0.11 ± 0.08
Crustose coralline algae		3.59 ± 2.41	13.43 ± 3.44	21.46 ± 3.67	18.45 ± 4.11	4.62 ± 1.85	3.13 ± 0.93
<i>Gelidium arbuscula</i>		3.57 ± 1.71	5.22 ± 1.93	5.54 ± 1.35	2.36 ± 1.72	5.71 ± 1.88	2.05 ± 1.43
<i>Gelidium canariense</i>		23.19 ± 5.13	1.68 ± 0.85	27.57 ± 5.66	-	3.01 ± 1.40	10.42 ± 4.23
<i>Grateloupia imbricata</i>		-	-	7.11 ± 1.87	-	-	0.09 ± 0.09
<i>Laurencia</i> sp.		0.05 ± 0.05	2.89 ± 1.14	0.07 ± 0.70	-	5.12 ± 2.67	0.69 ± 0.32
<i>Pterocladia capillacea</i>		69.06 ± 4.48	58.01 ± 6.23	5.36 ± 1.82	58.81 ± 6.30	38.92 ± 6.13	59.53 ± 5.49
<i>Rhodomenia</i> sp.		0.11 ± 0.11	0.28 ± 0.28	-	-	0.04 ± 0.04	0.95 ± 0.66
Phaeophyceae							
<i>Gongolaria abies-marina</i>		-	-	0.26 ± 0.26	-	-	-
<i>Sargassum</i> sp.		-	-	1.71 ± 1.71	-	-	-
<i>Colpomenia sinuosa</i>		-	-	0.03 ± 0.03	-	-	-
Chlorophyta							
<i>Chaetomorpha</i> sp.	-	0.12 ± 0.06	-	0.002 ± 0.002	0.01 ± 0.01	0.01 ± 0.01	

Table 4 (continued)

Species	Buenavista	Los Silos	Garachico	El Guincho	San Juan	Puerto Cruz
<i>Codium</i> sp.	-	5.77 ± 1.67	0.19 ± 0.16	-	0.01 ± 0.01	-
<i>Ulva</i> sp.	-	0.07 ± 0.03	0.18 ± 0.10	0.02 ± 0.01	0.08 ± 0.05	0.04 ± 0.02
<i>Valonia utricularis</i>	-	-	-	-	0.12 ± 0.10	0.01 ± 0.01
Cyanobacteria	0.01 ± 0.01	7.18 ± 1.99	12.51 ± 3.89	9.71 ± 2.96	2.95 ± 1.55	4.50 ± 2.11
Turf	0.02 ± 0.02	2.25 ± 0.82	6.36 ± 2.87	3.61 ± 1.02	18.04 ± 3.52	11.78 ± 3.77

Changes in species distribution are a worldwide phenomenon under current global warming conditions. The southern limits of several fucalean species, such as *Himanthalia elongata*, *Fucus serratus*, and *F. vesiculosus*, have already shifted to cooler latitudes due to the increase in SST (Duarte et al. 2013; Nicastro et al. 2013). The endemic continental Mediterranean species *Carpodesmia amentacea* and *C. zosteroides* have exhibited

population reductions and even local extinctions (Thibaut et al. 2005; Buonomo et al. 2018), and the large fucoid *Scytothalia dorycarpa*, endemic in the temperate regions of southern Australia, experienced poleward contraction after a heat wave event (Smale and Wernberg 2013). Similarly, the distribution of *G. canariense* has been reduced by more than 90% in only 30 years in Tenerife, and observational recordings suggest the same for the

Fig. 3 Average percentage of cover (% ± SE) of the species that contributed more than 60% to the spatial (six localities) and temporal (three years) variability of the macroalgae assemblages according to the SIMPER analysis. CCA, coralline crustose algae; Turf, turf-forming algae

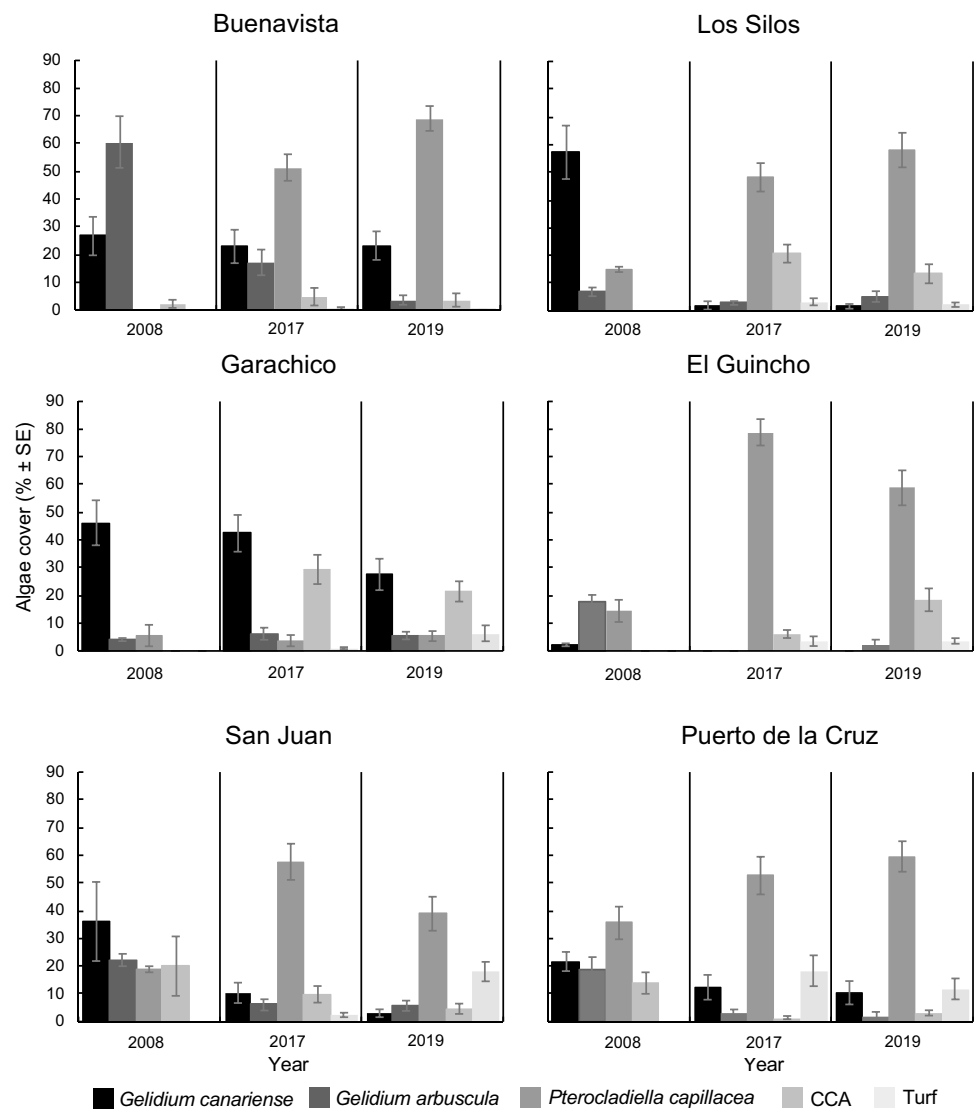
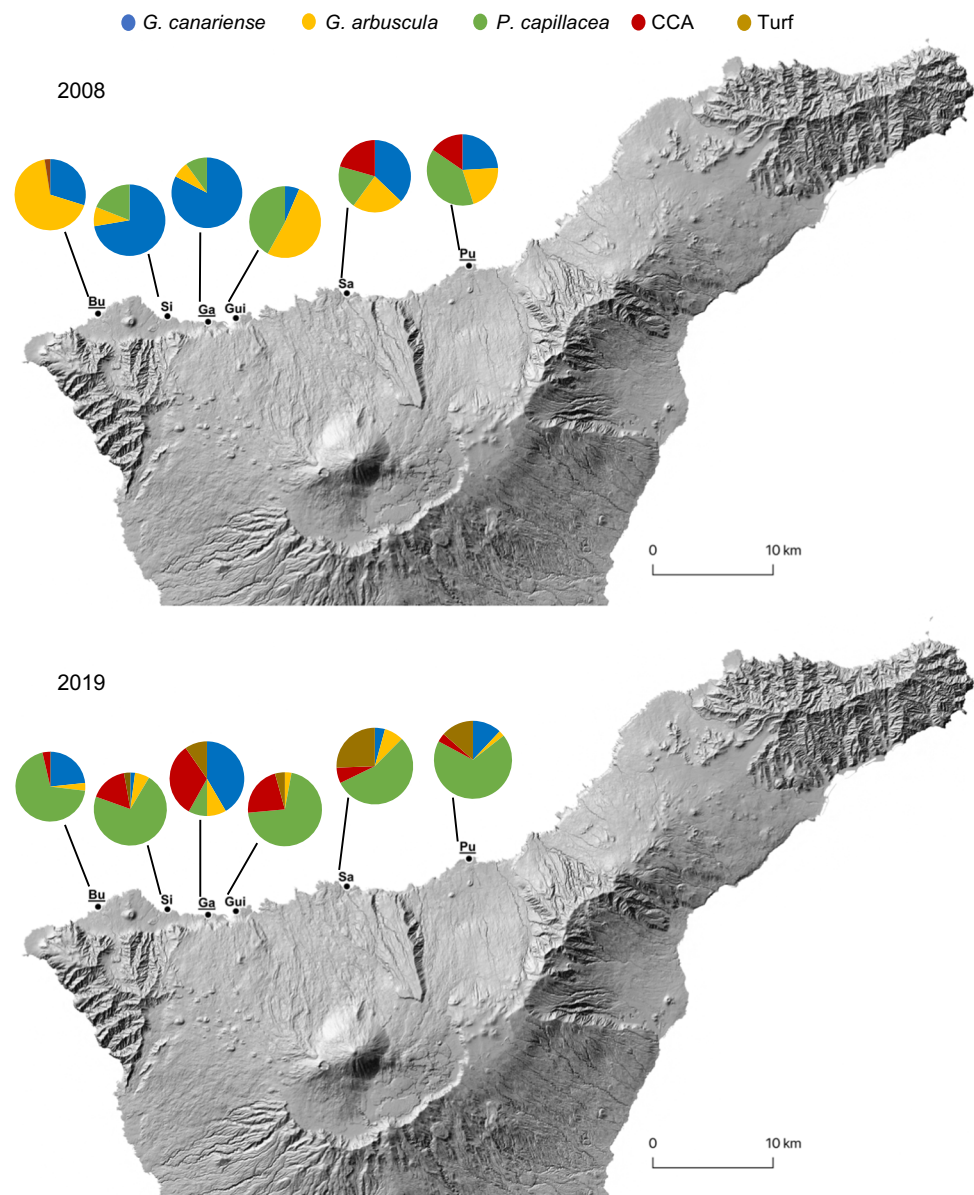


Fig. 4 Comparison between 2008 and 2019 of the average percentage of cover of the species that contributed more than 60% to the spatial and temporal variability of the macroalgae assemblages according to the SIMPER analysis (*Gelidium canariense*, *Gelidium arbuscula*, *Pterocladia capillacea*, coralline crustose algae (CCA) and turf-forming species) at all sites



other islands. Currently, the greatest populations of this species, at the westernmost site of the island, represent only 45% of its former distribution. However, contrary to endemic continental macroalgae, it is unfeasible for this species to migrate. On the one hand, it is impossible for propagules to drift northward, where other Macaronesian archipelagos are located, against the Canary current. On the other hand, the temperature of the African upwelling waters is markedly lower than that of Canarian waters, and the sediments that predominate along the coasts of the African upwelling complicate the settlement of propagules. Thus, as having a narrow geographic range is one

of the greatest extinction risk factors for many species (Lambers 2015), it is a matter of time before localized impacts or environmental changes, such as increasing temperature or UVR, cause the global extinction of this habitat-forming endemic macroalga from these oceanic islands.

Additionally, significant shifts in the presence or abundance of a habitat-forming macroalga species may lead to indirect effects on the associated benthic organisms. Different kinds of studies, including those with experimental conditions (Schiel and Lilley, 2011; Flukes et al. 2014; Álvarez-Losada et al. 2020), extreme

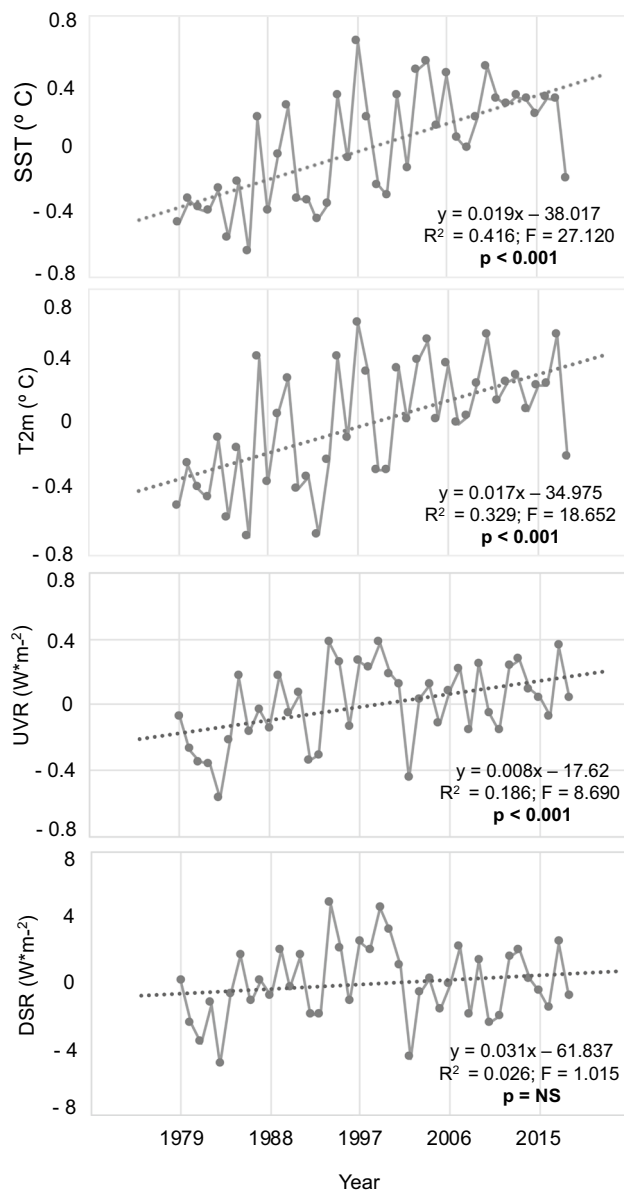


Fig. 5 Linear regression of SST (sea surface temperature), T2m (temperature 2 m above sea surface), UVR (downward ultraviolet radiation) and DSR (downward solar radiation) anomalies with time for the period 1979–2018. The linear trend equation, R^2 , F -statistic and p values are provided for each variable. NS, no significant

events (Smale and Wernberg, 2013; Borja et al. 2018), and continuous environmental changes (Díez et al. 2012; Tanaka et al. 2012), have shown that warming favors turf-forming algae over habitat-forming species. In southern Australia, the significant reduction in *S. dorycarpa* after an extreme heat wave occasioned

deleterious effects on the encrusting coralline algae and sponges, which were favored in the canopy, while turf-forming species proliferated significantly (Smale and Wernberg 2013). In *G. canariense* assemblages, we detected that over the last 10 years, the cover of CCA and turf-forming species has increased. We would like to point out here that in *G. canariense* assemblages, CCA are foundation species (Pinedo and Afonso-Carrillo 1994; Muñoz et al. 2007); therefore, CCA cover may have been underestimated in previous studies, when *G. canariense* assemblages were bushy. On the other hand, turf-forming species are considered tolerant to certain levels of environmental changes (Bertolini 2019), as highlighted in previous paragraphs.

The reduction in the dominant species (*G. canariense* and *G. arbuscula*) in Tenerife has mainly resulted in a free niche that is occupied by the cosmopolitan red alga *Pterocladia capillacea*, currently the most abundant species in most of the northern exposed coasts of the islands. It is important to indicate that the positive relationship of *P. capillacea* with SST, T2m, and UVR is markedly steep despite the absence of significance. This may be due to the similar cover data obtained in the 2017 and 2019 surveys, which masked the real relationship with the climatic variables. Nevertheless, this generalist species exhibits fast growth and high reproductive potential under a wide range of environmental conditions (Guiry and Guiry 2020), which makes it a strong competitor against *Gelidium* species.

In conclusion, the decline observed in assemblages of *G. canariense* and *G. arbuscula* is negatively correlated to environmental variables related to climate change. An increase in SST and T2m of 0.2 °C per decade is significantly correlated with the regression of the cover of both species. Consequently, this regression has left their niche available for *P. capillacea* to proliferate. Moreover, the rise in UVR is negatively correlated with the cover of *G. canariense* over the last decade. Although experimental studies are needed to determine the direct effects of these climatic variables on the species, the results of this study show the susceptibility of both species of *Gelidium* to global warming and especially highlight the disadvantage of *G. canariense*, revealing the need to upgrade the conservation status of this species. Understanding the spatiotemporal population trends of endemic macroalgae from oceanic islands, which grow and reproduce within narrow environmental and spatial boundaries, is crucial for predicting the future of many other species affected by the current climatic scenario, and it leads to an urgent call for marine biodiversity managers to implement conservation measures of these habitat-forming species.

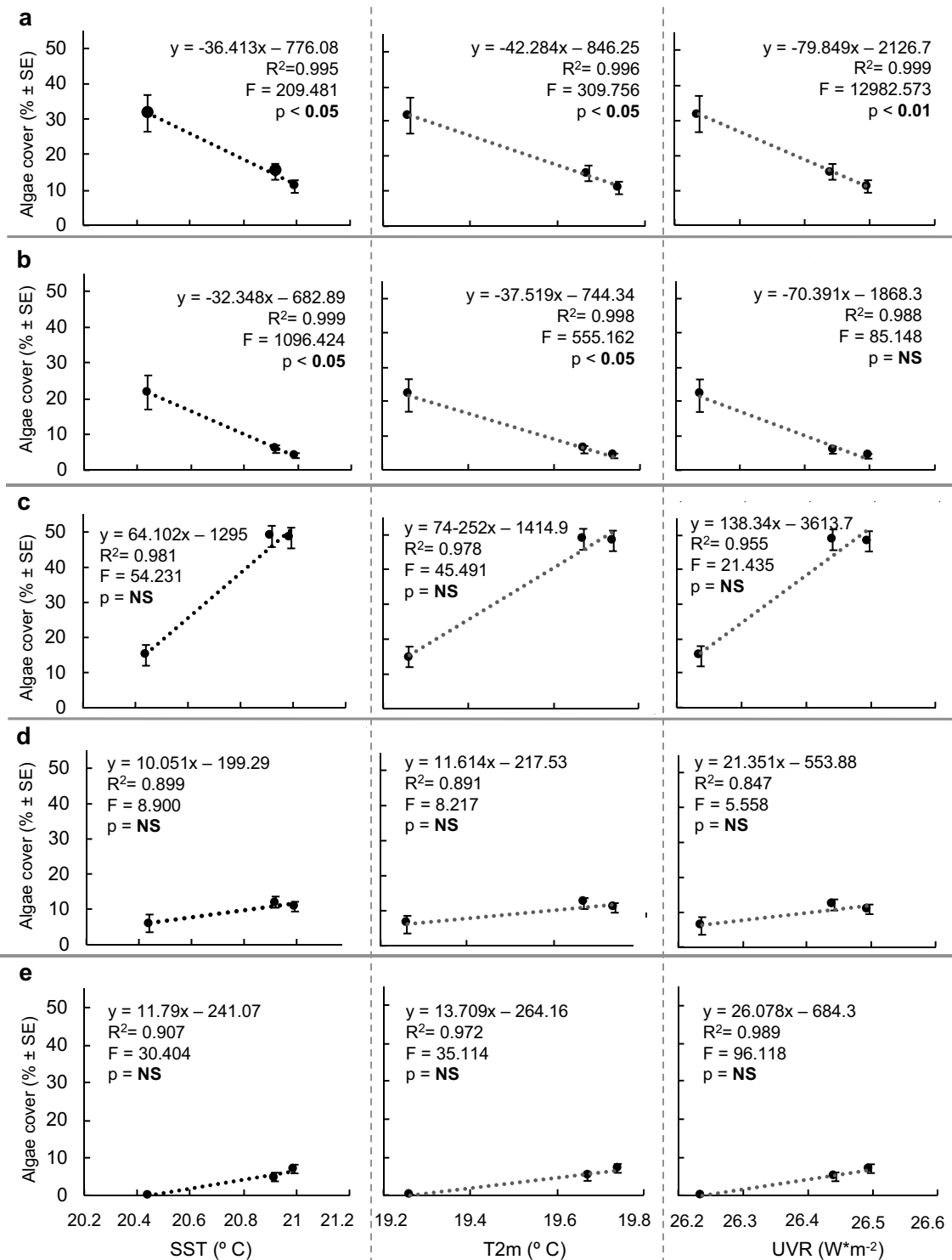


Fig. 6 Correlation of the percentage of cover of the species that contributed more than 60% to the variability of the macroalgae assemblages among sites and years: **A** *Gelidium canariense*, **B** *G. arbuscula*, **C** *Pterocliadiella capillacea*, **D** coralline crustose algae (CCA)

and **E** turf-forming species, with the climatic variables with significant increment for the period 1979–2018 (SST, sea surface temperature; T2m, temperature 2 m above sea surface; UVR, downward ultra-violet radiation). NS, no significant

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Declarations

Conflict of interest The authors declare no competing interests.

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