

EFFECT OF ULTRAVIOLET-B IRRADIATION ON FATTY ACIDS, AMINO ACIDS, PROTEIN CONTENTS, ENZYME ACTIVITIES AND ULTRASTRUCTURE OF SOME ALGAE

Nadia H. Noaman¹, Faiza M.A. Akl², Magda A. Shafik²,
Mohamed S.M. Abdel-Kareem¹, Wafaa M. Menesi²

¹Department of Botany, Faculty of Science, Alexandria University, Egypt

²Department of Biological and Geological Sciences, Faculty of Education,
Alexandria University, Egypt

e-mail: myabdelkareem@yahoo.com, msabdelkareem@gmail.com

Abstract. A series of experiments were conducted to determine fatty acids, amino acids, protein contents and enzymes activities of three algae (*Ulva lactuca*, *Sargassum hornschurchii* and *Pterocladia capillacea*) subjected to UV-B radiation. These parameters were estimated, when the UV-absorbing compounds contents recorded its maximum after the third day of irradiation of 60 minutes daily. Total saturated, mono unsaturated and polyunsaturated fatty acids of *U. lactuca* and *S. hornschurchii* were increased due to UV-B irradiation, while that of *Pterocladia capillacea* decreased. The study shows that amino acids of the three studied algal species varied greatly due to UV-B irradiation. The total protein content showed also different responses after UV-B irradiation. Exposure to UV-B radiation increased the activity of superoxide dismutase, ascorbate peroxidase and catalase of the three irradiated algal species. The findings also suggests that exposure to UV-B irradiance also affect the ultrastructure of all the irradiated algal species.

Keywords: Algae, ultraviolet, fatty acids, amino acids, proteins, enzyme, ultrastructure.

1. Introduction

The ozone layer is vital to life on Earth because it is the principal agent that absorbs ultraviolet radiation (UVR) in the Earth's atmosphere. Over the last 50 years, stratospheric ozone has decreased about 5% (Pyle, 1997). Depletion of ozone layer is due to anthropogenically released atmospheric pollutants such as chlorofluorocarbons (CFC_s), chlorocarbons and organobromides (Weatherhead & Andersen, 2006).

Detection of the reduction of stratospheric ozone layer led to the interest in investigating the effects of UV-B radiation on algae since radiation has the ability of reaching algae at any position due to the fact that radiation are capable even of penetrating the water column to significant depth (Tian & Yu, 2009).

The damaging effect of UV on algae include growth (Liu *et al.*, 2007), morphology (Wu *et al.*, 2005), reproduction (Cordiet *et al.*, 2001), motility (Häder, 1993), respiration (Kashian *et al.*, 2004), photobleaching of chlorophyll a, reduced photosynthesis (Rautenberger & Bischof, 2008), reduction of chlorophyll content (Zverdanovic *et al.*, 2009), degradation of light harvesting proteins (Jr, 2005), changing of protein profile (Abdel-Kareem, 1999), inhibition of enzymes of

nitrogen metabolism (Sinha&Häder, 2000) and other enzyme activities (Shiu and Lee, 2005).

However, there are few published papers pertaining to the effects of UVR on the fatty acids composition of microalgae (e.g. Wang & Chai, 1994 and Odmark *et al.*, 1998) and those that have been published are seemingly contradictory. The extant literature on the subject includes: some studies reporting an overall increase in SFA (saturated fatty acids) and MUFA (monounsaturated fatty acids) and decrease in PUFA (polyunsaturated fatty acids) upon UVR (e.g. Wang & Chai, 1994), two papers reported no significant differences in the fatty acid profile between UVBR treatments and the control (Skerratt *et al.*, 1998), and other reported an increase in UFA after UVR exposure (Gupta *et al.*, 2008).

Reactive oxygen species (ROS), such as hydroxyl ions, superoxide anions, and peroxy radicals, are involved in oxidative damage to cell components, regulation of signal transduction and gene expression, and inactivation of receptors and nuclear transcription factors when overproduced (Imlay & Linn, 1998). Subsequently it leads to many clinical diseases due to oxidative stress provided by these kinds of free radicals.

The response of superoxide dismutase (SOD), the first line of defence against ROS in plants (Alscher *et al.*, 2002), to UV stress depends on the algal species and the exposure time. SOD activity in the green macro alga *Ulva fasciata* increased by UV radiation (Shiu and Lee, 2005) and in the red macroalga *Corallina officinalis* (Li *et al.*, 2010). The increase of SOD and catalase (CAT) activities by UV radiation were greater in *Gelidiummamsii* than *Pterocladia capillacea*. UV radiation also increased SOD activity in symbiotic dinoflagellate (Lesser and Shick, 1989) but long term UV exposure decreased SOD activity in *Chlorella vulgaris* (Malanga *et al.*, 1997). Macroalgae (*Monostromaarcticum*, *Acrosiphoniapenicilliformis*, *Coccotylus truncates*, *Phycodrysrubens*, *Palmaria palmate* and *Devaleraearamentacea*) show less SOD induction by UV and it is even depressed in some species (Aguilera *et al.*, 2002). The induction of (CAT) by UV radiation was also observed in algae (Levy *et al.*, 2006). Because of the low affinity of CAT for H₂O₂ but the high affinity of APX (Ascorbate peroxidase) for H₂O₂, the higher induction of CAT activity accompanied by depression of APX activity & higher H₂O₂ production in *G. Amnsi* seems to indicate that this species faces greater oxidative stress upon exposure to UV radiation than *Pterocladia capillacea* (Lee and Shui, 2009).

A number of algae that are simultaneously exposed to visible and UV radiation have evolved mechanisms as accumulation of detoxifying enzymes and antioxidants (Mittler and Tel-Or, 1991) and synthesis of UV-protectants (Oren and Gunde-Cimerman, 2007).

In radiation exposure experiments, the effects of mild artificial UV conditions on ultrastructure of two red algal species *Palmariapalmata* and *Odonthaliadentata* from the Arctic have been investigated. The transmission electron microscope (TEM) results demonstrated that the photosynthetic apparatus was severely influenced by UV in both species (Holzinger *et al.*, 2004). Also the green alga *Dunaliellasalina* had many changes in ultrastructures during acclimation to enhanced UV-B radiation (Tian and Yu, 2009).

The present study aimed to study the defense mechanism(s) created by some algal cells against the effect of harmful UV-B radiation. For this purpose the influence of UV-B radiation on amino acids, fatty acids, protein contents enzyme activities and the ultrastructure of the selected algae has been studied.

2. Materials and methods

Algal species: *Ulva lactuca*, *Sargassum hornschurchii* and *Pterocladia capillacea* were collected in late July 2009 from Abu Qir in Alexandria. After harvesting, whole algae were extensively washed several times with natural sea water to remove any attached sand and the rhizoidal portions were removed to avoid microbial contamination. Then the algal materials were conveyed to the laboratory in plastic bags filled with sea water.

Culturing conditions and UV-irradiation: The whole algae were rinsed and placed in shallow trays with aeration inlet. Water used for culturing was collected from the sampling site. The trays were placed in an environmental cabinet at $30\pm 2^\circ\text{C}$ for 24 hours. In irradiation experiment, the samples were placed in Petri dishes (15 cm diameter) without covers and exposed directly to UV light. The source of light was UV VL-8.LM lamp supplied by VILBER LOURMAT-France. The UV-B light intensity of the lamp at distance 15 cm was $660 \mu\text{W cm}^{-2}$ and the supplied light has the wave length 312 nm. Samples were irradiated for 20, 40 & 60 minutes daily for 5 days at distance 15 cm.

Fatty acids determination: It was performed according to Radwan (1978).

Amino acids determination: Amino acid determination was performed according to the method of Winder and Eggum (1966). The system used for the analysis was high performance, amino acid analyzer, (SYKAM Amino acids analyzer, version 6.8).

Protein determination: Five days irradiated samples were used for total proteins determination according to Hartree (1972).

Enzyme activities estimation: Superoxide oxide dismutase was performed according to Giannopolitis and Ries (1977), ascorbate peroxidase according to Nakano and Asada (1981), while catalase activity was measured as described by Beers and Sizer (1952).

Ultrastructure: It was performed according to Reynolds (1963) and Mercer & Birbeck (1966). Then examined by Philips 400 T electron microscope at 60 – 80 KV.

Statistical analysis: The effect of UV-B radiation on fatty acids, amino acids contents were evaluated by means of paired samples t-test on the parameters estimated before and after exposure to UV-B radiation using SPSS version 10.0. Differences were considered significant at $P \leq 0.05$. The effect of UV-B radiation on these parameters was considered to be statistically significant at a level of $P < 0.05$.

3. Results

1. Fatty acids contents

Table (1) indicated that all fatty acids of *Ulva lactuca* increased after exposure to UV-B radiation for three days (60 minutes daily) except the saturated fatty acids C12:0 and C15:0, the monounsaturated fatty acid C14:1 and the polyunsaturated fatty acid C18:2, which decreased by 0.162, 0.325, 0.421 and 0.172 $\mu\text{g g}^{-1}$ fresh weight, respectively. Four fatty acids (C20:0, C17:1, C20:5 and C22:2) appeared in the irradiated alga, which were completely absent in the control.

The most affected fatty acids by UV-B irradiation was C14:0, which increased forty times compared to control, also C13:0, C15:1 and C22:6, which were increased by nearly four folds their initial concentration of the control. The less affected fatty acids of *U. lactuca* by UV-B irradiation were C8:0, C10:0 and C17:0, which increased by 0.017, 0.022 and 0.077 $\mu\text{g g}^{-1}$ fresh weight), respectively.

Table 1. Fatty acids content of *Ulva lactuca* before (control) and after (irradiated) exposure to UV-B radiation of 60 minutes daily for three days.

Fatty acid	Fatty acid content ($\mu\text{g g}^{-1}$ fresh weight)			
	Control	Irradiated	Increase/decrease	
Saturated fatty acids	C6:0	0.054	0.182	0.128
	C8:0	0.016	0.033	0.017
	C10:0	0.008	0.030	0.022
	C12:0	0.196	0.034	-0.162
	C13:0	0.811	2.896	2.085
	C14:0	0.046	1.763	1.717
	C15:0	0.406	0.081	-0.325
	C16:0	0.927	2.227	1.3
	C17:0	0.042	0.119	0.077
	C18:0	0.073	0.194	0.121
C20:0	-	0.009	0.009	
Total	2.579	7.568	4.989	
Mono unsaturated fatty acids	C14:1	0.474	0.053	-0.421
	C15:1	0.480	1.729	1.249
	C16:1	0.232	0.828	0.596
	C17:1	-	0.025	0.025
	C18:1	0.547	0.744	0.197
	C22:1	0.070	0.206	0.136
Total	1.803	3.585	1.782	
Poly unsaturated fatty acids	C18:2	0.262	0.090	-0.172
	C18:3	0.113	0.249	0.136
	C20:3	0.053	0.445	0.392
	C20:5	-	0.038	0.038
	C22:2	-	0.078	0.078
C22:6	0.544	1.973	1.429	
Total	0.972	2.873	1.901	
Total fatty acids	5.354	14.026*	8.672	

(*) Marked differences are significant at $p \leq 0.05$

The data in Table 1 showed that the total saturated and polyunsaturated fatty acids of *U. lactuca* increased approximately three times after exposure to UV-B radiation, while monounsaturated fatty acids increased nearly two times their initial contents of the control. Total fatty acids content in irradiated *U. lactuca* increased approximately three folds comparing to control.

Table 2. Fatty acids content of *Sargassum hornschurchii* before (control) and after (irradiated) exposure to UV-B radiation of 60 minutes daily for three days.

Faty acid	Fatty acid content ($\mu\text{g g}^{-1}$ fresh weight)			
	Control	Irradiated	Increase/decrease	
Saturated fatty acids	C6:0	0.014	0.095	0.081
	C8:0	0.016	0.071	0.055
	C10:0	-	0.015	0.015
	C11:0	0.022	0.017	-0.005
	C12:0	0.090	0.249	0.159
	C13:0	0.497	0.021	-0.476
	C14:0	0.137	0.357	0.22
	C15:0	0.031	0.611	0.58
	C16:0	1.351	3.292	1.941
	C17:0	0.011	0.009	-0.002
	C18:0	0.037	0.135	0.098
	C20:0	0.105	0.344	0.239
	C21:0	-	0.033	0.033
C23:0	0.020	0.029	0.009	
	Total	2.331	5.279	2.948
Mono unsaturated fatty acids	C14:1	0.332	0.704	0.372
	C15:1	0.382	0.759	0.337
	C16:1	0.176	0.308	0.132
	C17:1	0.008	0.035	0.027
	C18:1	0.052	-	0.052
	C20:1	0.014	0.044	0.03
	C22:1	0.071	0.047	-0.024
	Total	0.995	2.213	1.218
Poly unsaturated fatty acids	C18:2	0.009	0.013	0.004
	C18:3	0.008	0.064	0.056
	C20:2	0.065	0.145	0.08
	C20:3	0.031	0.154	0.123
	C20:5	0.010	0.022	0.012
C22:6	0.654	1.254	0.6	
	Total	0.777	1.652	0.223
Total fatty acids	4.103	8.828*	4.725	

(*) Marked differences are significant at $p \leq 0.05$

Results in Table 2 illustrated that all fatty acids of *Sargassum hornschurchii* increased after exposure to UV-B radiation for three days except the three saturated fatty acids (C11:0, C13:0 and C17:0), which decreased by 0.005, 0.476 and 0.002 $\mu\text{g g}^{-1}$ fresh weight, respectively and the monounsaturated fatty acid C22:1 which decreased by 0.024 $\mu\text{g g}^{-1}$ fresh weight. The most increased fatty acid of *S. hornschurchii*, due to UV- B irradiation was the saturated fatty acid C16:0, which increased by 1.941 $\mu\text{g g}^{-1}$ fresh weight. The less increased fatty acids were the saturated fatty acid C23:0, which increased by 0.009 $\mu\text{g g}^{-1}$

fresh weight and the polyunsaturated fatty acid C18:2, which increased by 0.004 $\mu\text{g g}^{-1}$ fresh weight.

It must be mentioned that the two saturated fatty acids, C10:0 and C21:0 appeared in the irradiated alga, which were completely absent in the control. Meanwhile the mono unsaturated fatty acid C18:1 was completely disappeared after exposure to UV-B radiation.

Table 2 showed that the contents of total saturated, mono unsaturated and polyunsaturated fatty acids of *S. hornschurchii* were doubled due to exposure to UV-B irradiation. Consequently, the total fatty acids content in irradiated alga increased also two times compared to the control. The data in Table 3 showed that all concentrations of fatty acids of *Pterocladia capillacea* decreased after exposure to UV-B radiation for three days except C13:0, C15:0, C14:1, C15:1, C16:1, C20:3 and C22:6 where they increased by 2.195, 0.114, 0.165, 0.3, 0.098, 0.002, 0.187 $\mu\text{g g}^{-1}$ fresh weight, respectively.

Table 3. Fatty acids content of *Pterocladia capillacea* before (control) and after (irradiated) exposure to UV-B radiation of 60 minutes daily for three days.

Fatty acid	Fatty acid content ($\mu\text{g g}^{-1}$ fresh weight)			
	Control	Irradiated	Increase/decrease	
Saturated fatty acids	C6:0	0.180	0.053	-0.127
	C8:0	0.017	0.014	-0.003
	C10:0	0.045	0.007	-0.038
	C11:0	0.032	0.022	-0.01
	C12:0	0.458	0.005	-0.453
	C13:0	0.039	2.234	2.195
	C14:0	0.813	0.356	-0.457
	C15:0	1.130	1.244	0.114
	C16:0	4.898	2.400	-2.490
	C17:0	0.159	0.079	-0.08
	C18:0	0.574	0.204	-0.37
	C20:0	0.144	-	-0.144
Total	8.512	6.744	-1.768	
Mono unsaturated fatty acids	C14:1	1.192	1.357	0.165
	C15:1	1.192	1.492	0.3
	C16:1	0.629	0.727	0.098
	C17:1	0.043	0.019	-0.024
	C18:1	2.084	1.002	-1.082
	C22:1	0.079	0.024	-0.055
Total	5.219	4.621	-0.598	
Poly unsaturated fatty acids	C18:2	0.443	0.086	-0.357
	C18:3	0.133	0.022	-0.111
	C20:2	0.125	-	-0.125
	C20:3	0.092	0.094	0.002
	C20:4	1.027	0.373	-0.654
	C20:5	0.357	0.165	-0.192
	C22:6	1.363	1.550	0.187
Total	3.540	2.290	-1.25	
Total fatty acids	17.271	13.655	-3.616	

The most increased fatty acid of *Pterocladia capillacea* due UV-B irradiation was C13:0, where it increased 57 times compared to control, while the less increased fatty acid was C20:3 where it increased by only 0.002 $\mu\text{g g}^{-1}$ fresh weight. The most decreased fatty acids due to UV-B irradiation were C16:0 and C18:1, where they decreased nearly to its half contents compared to control. It must be mentioned that two fatty acids of *P. capillacea* (C20:0 and C20:2) were completely disappeared after exposure to UV-B radiation.

Table 3 showed that the contents of total saturated, mono and polyunsaturated fatty acids of *Pterocladia capillacea* decreased after exposure to UV-B radiation by 1.768, 0.598 and 1.25 $\mu\text{g g}^{-1}$ fresh weight, respectively. UV-B radiation caused the saturated fatty acids content to be dropped by 20.8% than the control. Mono unsaturated and polyunsaturated fatty acids in irradiated samples were found to be less than control by 11.5% and 35.3%, respectively.

It is noteworthy to mention that the increase of fatty acid contents due to UV-B irradiation in both *Ulva lactuca* and *Sargassum hornschurchii* was statistically significant at $P \leq 0.05$, meanwhile the decrease in this content in *Pterocladia capillacea* was statistically insignificant at $P \leq 0.05$.

2. Amino acids contents

Table 4 illustrated the contents of amino acids groups of *Ulva lactuca* before and after exposure to UV-B radiation for 60 minutes daily for three days. It was noticed that half of amino acids of *Ulva lactuca* increased after exposure to UV-B radiation for three days, while the other half of amino acids was decreased. The most increased amino acid was the aliphatic amino acid alanine, where it increased by 1.243 mg g^{-1} fresh weight and the less increased amino acid was valine, where it increased by 0.210 mg g^{-1} fresh weight.

The most decreased amino acid was the basic amino acid arginine, where it decreased by 4.468 mg g^{-1} fresh weight and the less decreased amino acid was the aliphatic amino acid leucine, which decreased by 0.013 mg g^{-1} fresh weight. Table 4 showed that the total acidic, aliphatic, aromatic and secondary amino acids groups of *U. lactuca* increased by 0.478, 50.681, 0.563 and 0.276 mg g^{-1} fresh weight, respectively. The most increased amino acid group was the aliphatic amino acid, where it increased nearly three folds their initial concentration of the control, but basic and sulphur-containing amino acids groups decreased by 6.355 and 0.064 mg g^{-1} fresh weight, respectively. Total amino acids content of *U. lactuca* decreased by 5.07 mg g^{-1} fresh weight after exposure to UV-B radiation for 60 minutes for three days.

Amino acids contents of *Sargassum hornschurchii* were presented in Table (5). It was noticed that all amino acids contents decreased after exposure to UV-B radiation for three days except the two basic amino acids histidine and lysine, and the aliphatic amino acid serine, which increased by 0.257, 0.232 and 0.989 mg g^{-1} fresh weight, respectively.

The most increased amino acid of *S. hornschurchii* due to UV-B irradiation was serine, where it increased by 0.989 mg g^{-1} fresh weight and the less increased was lysine where it increased by 0.232 mg g^{-1} fresh weight. The most decreased

amino acids by UV-B irradiation was the acidic amino acid glutamic where decreased by 9.34 mg g⁻¹ fresh weight and the less decreased one was the aromatic amino acid tyrosine, where it decreased by 0.187 mg g⁻¹ fresh weight.

Table 4. Amino acids groups content of *Ulva lactuca* before (control) after (irradiated) exposure to UV-B radiation of 60 minutes daily for three days.

Amino acid		Amino acid content (mg g ⁻¹ fresh weight)		
		Control	Irradiated	Increase/decrease
Acidic amino acids	Aspartic	13.935	13.887	-0.048
	Glutamic	14.681	15.207	0.526
	Total	28.616	29.094	0.478
Basic amino acids	Arginine	6.539	2.071	-4.468
	Histidine	3.130	1.842	-1.280
	Lysine	1.234	0.635	-0.597
	Total	10.903	4.548	-6.355
Aliphatic amino acids	Threonine	6.635	7.241	0.606
	Serine	16.206	12.941	-3.265
	Glycine	12.922	13.715	0.793
	Alanine	13.834	15.077	1.243
	Valine	9.146	9.356	0.21
	Leucine	10.074	10.061	-0.013
	Total	24.730	75.411	50.681
Aromatic amino acids	Tyrosine	0.407	0.070	-0.337
	Phenyl alanine	6.513	7.413	0.9
	Total	6.920	7.483	0.563
Sulphur -containing amino acids	Methionine	1.398	1.334	-0.064
Secondary amino acids	Proline	9.607	9.883	0.276
Total		132.825	127.753	-5.072

Table 5 represent the contents of amino acids groups of *S. hornschurchii* before and after exposure to UV-B radiation for 60 minutes daily for three days. These results showed that all total acidic, basic, aliphatic, aromatic, secondary and sulphur-containing amino acids decreased after exposure to UV-B radiation by 11.205, 0.032, 4.424, 0.704, 1.361 and 0.567 mg g⁻¹ fresh weight, respectively. All amino acids of *Pterocladia capillacea* increased after exposure to UV-B radiation for three days except the aliphatic amino acid serine and the aromatic amino acid tyrosine, where they decreased by 13.887 and 0.02 mg g⁻¹ fresh weight, respectively (Table 6).

The highest increase in amino acids content of *P. capillacea* due to UV-B irradiation was the aliphatic amino acid isoleucine, where it increased by 13.65 mg g⁻¹ fresh weight, while the less increased one was the sulphur-containing amino acid methionine where it increased only by 0.299 mg g⁻¹ fresh weight. The most decreased amino acid due to UV-B irradiation was the aliphatic amino acid serine (13.887 mg g⁻¹ fresh weight) and the less decreased amino acid was tyrosine (0.02 mg g⁻¹ fresh weight).

Table 5. Amino acids groups content of *Sargassum hornschurchii* before (control) and after (irradiated) exposure to UV-B radiation of 60 minutes daily for three days.

Amino acid	Amino acid content (mg g ⁻¹ fresh weight)			
	Control	Irradiated	Increase/decrease	
Acidic amino acids	Aspartic	9.764	7.899	-1.865
	Glutamic	16.178	6.838	-9.34
	Total	25.942	14.737	-11.205
Basic amino acids	Arginine	2.651	2.130	-0.521
	Histidine	0.576	0.833	0.257
	Lysine	1.039	1.271	0.232
	Total	4.266	4.234	-0.032
Aliphatic amino acids	Threonine	3.436	2.113	-1.323
	Serine	4.327	5.316	0.989
	Glycine	5.351	4.907	-0.444
	Alanine	5.025	3.540	-1.485
	Valine	3.860	3.535	-0.325
	Leucine	5.596	4.321	-1.275
	Isoleucine	4.071	3.507	-0.564
Total	31.666	27.239	-4.427	
Aromatic amino acids	Tyrosine	0.236	0.049	-0.187
	Phenyl alanine	3.124	2.607	-0.517
	Total	3.360	2.656	-0.704
Sulphur –containing amino acids	Methionine	1.024	0.457	-0.567
Secondary amino acids	Proline	5.476	4.115	-1.361
Total	71.734	53.438	-18.296	

Table 6. Amino acids groups content of *Pterocladia capillacea* before (control) and after (irradiated) exposure to UV-B radiation of 60 minutes for three day.

Amino acid	Amino acid content (mg/g fresh weight)			
	Control	Irradiated	Increase/decrease	
Acidic amino acids	Aspartic	15.399	22.867	7.468
	Glutamic	17.202	28.460	11.258
	Total	32.601	51.327	18.726
Basic amino acids	Arginine	8.298	10.198	1.900
	Histidine	1.268	2.262	0.976
	Lysine	1.699	2.040	0.341
	Total	11.265	14.500	3.235
Aliphatic amino acids	Threonine	8.717	11.071	2.354
	Serine	20.945	7.058	-13.889
	Glycine	15.670	18.471	2.801
	Alanine	13.761	15.368	1.607
	Valine	12.153	15.790	3.637
	Leucine	13.444	16.671	3.227
	Isoleucine	11.682	25.332	13.650
Total	96.372	109.761	13.389	
Aromatic amino acids	Tyrosine	0.050	0.030	-0.02
	Phenyl alanine	8.190	10.669	2.479
	Total	8.240	10.699	2.459
Sulphur -containing amino acids	Methionine	2.344	2.643	0.299
Secondary amino acids	Proline	11.735	15.220	3.485
Total	162.56	204.150*	41.593	

All the total acidic, basic, aliphatic, aromatic, secondary and sulphur-containing amino acids groups increased by UV-B irradiation by 18.726, 3.235, 13.389, 2.459, 3.485 and 0.299 mg g⁻¹ fresh weight, respectively. The most increased content was noticed in the acidic amino acids group where it increased by 57.4% compared to control. Total amino acids content of *P. capillacea* increased by 41.593 mg g⁻¹ fresh weight (25.6%) after exposure to UV-B radiation (Table 6).

It was found from the statistical analysis that there are no significant differences at $P \leq 0.05$ between the mean value of the total amino acids contents of the both *U. lactuca* and *S. hornschurchii* before and after treatments by UV-B radiation, while this mean values was statistically significant in the case of *P. capillacea* at $P \leq 0.05$ (Appendix 16).

3. Protein contents

Protein contents were determined in the samples of the three algae (*Ulva lactuca*, *Sargassum hornschurchii* and *Pterocladia capillacea*), which contained maximum UV-absorbing compounds. Table 7 indicated that the total protein content increased in *U. lactuca* through out the irradiation experiment. This increase was notably large after the first dose of irradiation, where its value was 11.20 mg g⁻¹ fresh weight and then the increment was gradually quite little by increasing the irradiation dose, where its values were 11.83, 12.41, 12.43 & 12.64 mg g⁻¹ fresh weight, respectively.

The protein contents of irradiated *S. hornschurchii* increased gradually through out the irradiation experiment. This increase was large after the first and second doses of irradiation, where their values were 16.35 & 17.67 mg g⁻¹ fresh weight, respectively. The increments were gradually quite little by increasing the irradiation dose, where their values were 18.00, 18.19 & 18.20 mg g⁻¹ fresh weight, respectively. *P. capillacea* showed notable decreases of protein contents after the first and second doses, where their values were 22.72 & 21.46 mg g⁻¹ fresh weight, respectively and also a gradual quite little decrease by increasing the following three irradiation doses, where their values were 21.04, 20.72 & 20.71 mg g⁻¹ fresh weight.

Table 7. Total protein content of *Ulva lactuca*, *Sargassum hornschurchii* and *Pterocladia capillacea* before (control) and after (irradiated) exposure to UV-B radiation of 60 minutes daily for five days.

Species	Content of protein (mg g ⁻¹ fresh weight).					
	Control	1 st day	2 nd day	3 rd day	4 th day	5 th day
<i>Ulva lactuca</i>	10.46	11.20	11.83	12.41	12.43	12.64
<i>Sargassum hornschurchii</i>	14.72	16.35	17.67	18.00	18.19	18.20
<i>Pterocladia capillacea</i>	24.72	22.72	21.46	21.04	20.72	20.71

4. Enzymes activities

Enzymes activities of the three algal species (*Ulva lactuca*, *Sargassum hornschurchii* and *Pterocladia capillacea*) were estimated after irradiation by UV-B for three days for 60 minutes. Table (8) showed that the activity of ascorbate peroxidase (APO) increased in all the three algae (*U. lactuca*, *S. hornschurchii* and *P. capillacea*) as a result of UV-B irradiation for three days by nearly 101.4, 25.2 and 43.5%, respectively. It was noticed that superoxide dismutase activity (SOD) in *U. lactuca* increased due to UV-B irradiation by approximately 51.2% compared to control. SOD activity in *S. hornschurchii* increased also by approximately 19.7%, while the highest increase (126.3%) of this activity was recorded by *P. capillacea* (Table 9). Catalase activity (CAT) was remarkably increased after UV-B irradiation (Table 10) in all the three algal species for three days. These increases were approximately 32.3, 56.4 and 35.9% for *U. lactuca*, *S. hornschurchii* and *P. capillacea*, respectively.

5. Ultrastructure of the algal species

The electron micrograph of *U. lactuca* before UV-B irradiation showed arrangement of the cell components and clear cell wall (CW) (Plate 1). When the cells were exposed to

Table 8. Ascorbate peroxidase activity of *Ulva lactuca*, *Sargassum hornschurchii* and *Pterocladia capillacea* before (control) and after (irradiated) exposure to UV-B radiation of 60 minutes daily for three days.

Species	Enzyme activity ($\mu\text{mol H}_2\text{O}_2 \text{ minutes}^{-1} \text{ g}^{-1}$ fresh weight)			
	Control	Treated	Increase/decrease	%Increase/decrease
<i>Ulva lactuca</i>	1.488	2.997	1.509	101.4
<i>Sargassum hornschurchii</i>	0.329	0.412	0.083	25.2
<i>Pterocladia capillacea</i>	0.411	0.590	0.179	43.5

Table 9. Superoxide dismutase activity of *Ulva lactuca*, *Sargassum hornschurchii* and *Pterocladia capillacea* before (control) and after (irradiated) exposure to UV-B radiation of 60 minutes daily for three days.

Species	Enzyme activity (unit gm^{-1} fresh weight)			
	Control	Irradiated	Increase/decrease	%Increase/decrease
<i>Ulva lactuca</i>	0.43	0.65	0.220	51.2
<i>Sargassum hornschurchii</i>	0.66	0.79	0.130	19.7
<i>Pterocladia capillacea</i>	0.38	0.86	0.480	126.3

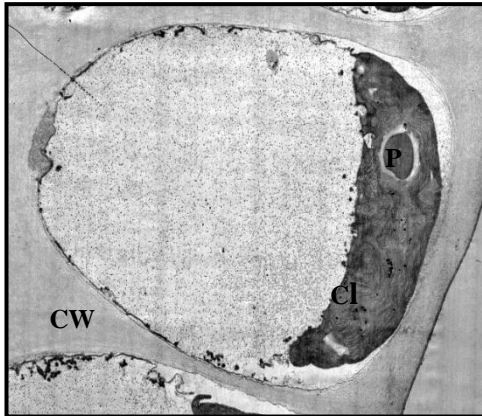


Plate (1): The electron micrograph of *Ulva lactuca* before UV-B irradiation showing chloroplasts (Ch), pyrenoid (P) and cell wall (CW) (2.5×10^3).

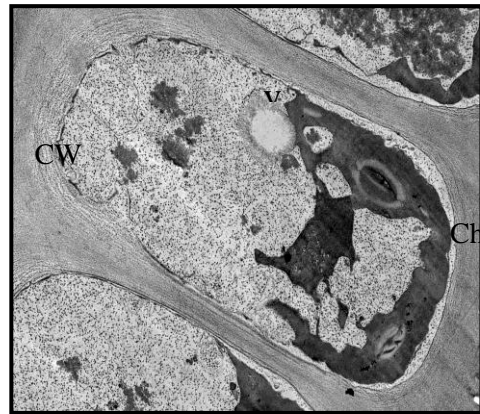


Plate (2): The electron micrograph of *Ulva lactuca* irradiated by UV-B of 60 minutes daily for five days showing dissipation of chloroplast (Ch) and appearance of a vacuole (V) and lamellated cell wall (CW) (2.5×10^3).

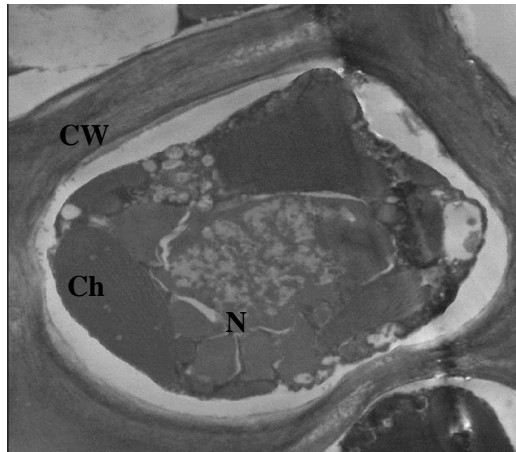


Plate (3): The electron micrograph of *Sargassum hornschurchii* before UV-B irradiation showing the typical chloroplast (Ch), cell wall (CW) with clear arrangement of thylakoids (7.5×10^3).

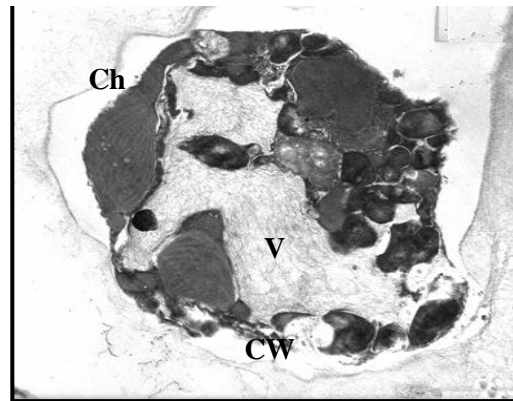


Plate (4): The electron micrograph of *Sargassum hornschurchii* irradiated by UV-B irradiation of 60 minutes daily for five days showing dramatic disorganization of cell components of the alga, irregularity of cell wall and appearance of vacuole (V) (7.5×10^3).

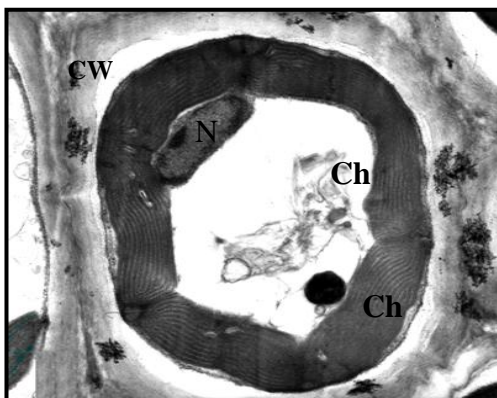


Plate (5): The electron micrograph of *Pterocladia capillacea* before UV-B irradiation showing the clear arrangement of thylakoids inside chloroplast (Ch), cell wall (CW) and nucleus (N).

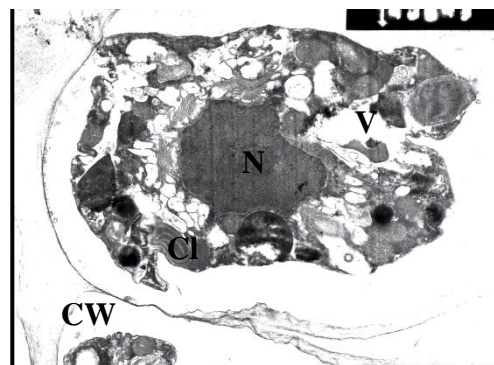


Plate (6): The electron micrograph of *Pterocladia capillacea* irradiated by UV-B irradiation of 60 minutes daily for five days showing disturbance of the cell inclusions, appearance of vacuoles (V), irregularity of cell wall (CW) and chloroplast structure (Ch) is less clear with disorganization of thylakoids.

Table 10. Catalase activity of *Ulva lactuca*, *Sargassum hornschurchii* and *Pterocladia capillacea* before (control) and after (irradiated) exposure to UV-B radiation of 60 minutes daily for three days.

Species	Enzyme activity ($\mu\text{mol H}_2\text{O}_2 \text{ minutes}^{-1} \text{ g}^{-1}$ fresh weight)			
	Control	Treated	Increase/decrease	% Increase/decrease
<i>Ulva lactuca</i>	26.894	35.576	8.682	32.3
<i>Sargassum hornschurchii</i>	11.647	18.211	6564	56.4
<i>Pterocladia capillacea</i>	13.552	18.423	4.871	35.9

60 minutes daily doses of UV-B radiation for five days, chloroplast showed dissipation and irregularity in shape and a small vacuole appeared (V). On the other hand, the cell wall and the pyrenoids appeared unaffected (Plate 2). The electron micrograph of untreated *S. hornschurchii* showed obvious arrangement of thylakoid membranes, obvious nucleus with visional nuclear envelope and cell wall (Plate 3). The irradiated cells showed some disorganization of cell components, malformation of the cell, wrinkled cell wall and appearance of some vacuols. The nucleus is not affected (Plate 4). Plate (5) showed untreated cell of *P. Capillacea* with typical chloroplasts that have obvious arrangement of thylakoids, well organized nucleus and visional cell wall. Meanwhile, irradiated cell (Plate 6) showed partially damaged cell wall, disturbance of the cell inclusions including chloroplasts and appearance of vacuoles.

6. Discussion

Algae have an important role as food for fish and crustaceans and their nutritional value is mainly related to the content of essential fatty acids. Some studies reporting an overall increase in saturated fatty acid and monounsaturated fatty acids and decrease in polyunsaturated fatty acids of algae (e.g. Wang & Chai, 1994), while others reported an increase in saturated fatty acid, monounsaturated & polyunsaturated fatty acids of algae (e. g. Noaman, 2007). Meanwhile, few papers reported no significant differences in the fatty acid profiles between UV radiation treatments and the control (e.g. Skerratt *et al.* 1998).

The composition of fatty acids of *Spirulina platensis* in response to UV-B radiation was found to have 23.5% saturated fatty acid (SFA), 76.4% monounsaturated fatty acid (MUFA) and polyunsaturated fatty acid (PUFA). In contrast to its UV-B untreated counterpart, SFA was 46.6% and MUFA and PUFA were 53.3%, which suggested that UV radiation reduces saturated fatty acid and increased unsaturated fatty acids in *S. platensis* (Gupta *et al.*, 2008). It is also observed that gamma linolenic acid was an important component of the total content of PUFAs of UV treated alga.

Liang *et al.* (2006) concluded that the effect of UV radiation on algal fatty acid compositions depends on algal species, the nitrogen concentration and time of UV radiation exposure. This was true for our results, since the total saturated,

mono- and polyunsaturated fatty acid contents of *U. lactuca* and *S. hornschurchii* (Tables 1 & 2) increased significantly due to UV-B irradiation, while these of *P. capillacea* were slight decreased (Table 3). Several studies showed that UV radiation resulted in an increase of (PUFA) and reduction of SFA. For example, Liang *et al.* (2006) who showed that UV radiation resulted in an increase of PUFA and reduction of SFA in the marine diatoms *Phaeodactylum tricornutum* and *Chaetoceros mulleri*, which agrees with the findings of De Lang and Van Donk (1997) for *Cryptomonas pyrenoidifera* and Skerratt *et al.* (1998) for *Phaeocystis antarctica*.

In the same topic, Meireles *et al.* (2003) recorded that UV irradiation increased the two fatty acids eicosapentaenoic and docosahexaenoic (n-3 fatty acids) by the alga *Pavlova lutheri*. Meanwhile there was a reduction of short-chained FA (C-14, C-16). UV increases fatty acids of *Chaetoceros simplex* (Boutry *et al.*, 1976) and *Pavlova lutheri* (Meireles *et al.*, 2003). Kobayashi (1998) found an increase in fatty acids of 18 carbon atoms by UV irradiation but that of 20-22 carbon atoms was not affected by the exposure time.

The increase of fatty acid contents of both *U. lactuca* and *S. hornschurchii* may be interpreted as a physiological response to adapt the alga to UV-B irradiation stress. Since PUFAs are known in regulating membrane fluidity (Hall *et al.*, 2002) and physiological processes under stress (Golecki and Drews, 1982). At the same time membrane lipid unsaturation increases tolerance of Cyanophyta to UV radiation (Ehling & Scherer, 1999). SFAs and MUFAs provide the energy required for rebuilding of the photosynthetic apparatus and PUFAs are essential for chlorophyll membrane development (Skerratt *et al.*, 1998). Meanwhile, the decrease of fatty acids content of *P. capillacea* may be due to splitting of fatty acids as a result of UV-B irradiation (Kobayashi, 1998) and/or lipid peroxidation (He *et al.*, 2002). The previous data are in complete agreement with the results of Goes *et al.*, (1994). They showed that the formation of PUFA in the green alga *Tetraselmis* sp. was suppressed by UV, the results of Skerratt *et al.* (1998) who reported that PUFA decreased in *Chaetoceros simplex* under UV radiation.

Noaman (2007) found that the drop in C18:3 in *Synechococcus leopoliensis* by its exposure to UV for 5 minutes was followed by the increase of that fatty acid of 18 carbon atoms with increasing the exposure time, while polyunsaturated fatty acid of 22 carbon atoms decreased by increasing the exposure time.

In contrast to Bhandari and Sharma (2006) who found that fatty acid profile of *Phormidium corium* did not show any qualitative changes due to exposure to UV-B irradiation. *Ulva lactuca* showed appearance of four new fatty acids (SFA C20:0, MUFA C17:1 and PUFAs C20:5 & C22:2), while *S. hornschurchii* showed appearance of two fatty acids (SFAs C10:0 & C21:0) and disappearance of one fatty acid (MUFA C18:0). At the same time two fatty acids (SFA C20:0 & PUFA C20:2) disappeared from the fatty acid profile of *P. capillacea*. Similar disappearance was reported by Noaman (2007) for *Synechococcus leopoliensis*, while UV radiation causes induction of PUFA C20:20 in the marine diatom *Chaetoceros simplex* (Boutry *et al.*, 1976). At the same time, UV effects on cell components (e.g. lipids, fatty acids, proteins, amino acids) and metabolic

processes have been studied (Karentz *et al.*, 1994). Meanwhile, UV irradiation may change the contents of proteins and amino acids of marine algae (Korbee *et al.*, 2005).

In the obtained literatures, we noticed no general or specific trend for the effect of UV-B radiation on the concentration of individual amino acids. Meanwhile, Döhler (1984) reported that the effect of UV-B radiation on concentration of amino acids was species-dependent. For example: some amino acids in *Synechococcus leopoliensis* as lysine and arginine decreased by the exposure to UV irradiation, while aspartic increased (Noaman, 2007). The same author showed that cysteine, alanine and valine completely disappeared from mutants M₁, M₂ and M₃. Exposure of *Scenedesmus quadricauda* to UV-A caused five (including proline) of 17 detected amino acids to increase, while only aspartic acid and histidine increased in UV-C treatment (Kovacik *et al.*, 2010). UV-A and UV-B irradiance resulted in an increase of main amino acid biosynthesis and an enhancement of the main free amino acids (Döhler *et al.*, 1997), results are discussed in relation to the UV-effects on photosynthetic pigments and the key enzymes of the carbon and nitrogen metabolism. This conclusion was noticed in our results, where some amino acids decreased and others increased due to UV-B irradiation in the three studied species (Tables 4-6).

The aromatic amino acid phenyl alanine, which can absorb UV-B radiation (Martin *et al.*, 1985) was found to increase in *U. lactuca* and *P. capillacea* (Tables 4 & 6). Meanwhile, the decrease of this amino acid in *S. hornschurchii* (Table 5) may be due to the formation of phenyl alanine ammonia-lyase in response to UV radiation (Campos *et al.*, 1991).

UV radiation accumulates proline that can protect plant cell against UV radiation induced peroxidative processes (Sarkar *et al.*, 2011). This was true for *U. lactuca* and *P. capillacea* in which proline increased due to UV-B irradiation (Tables 4 & 6). Since proline is one of the important solutes, which accumulate in many organisms by exposure to environmental stresses, it is likely that proline accumulation is related to the protection of these organisms against singlet oxygen production during stress conditions (Alia *et al.*, 1995). At the same time, proline accumulation may be critical for stimulating the pentose phosphate pathway in order to provide key precursors for the phenylpropanoid pathway (Kwok & Shetty, 1997). With the same respect, a three-fold increase in proline occurred in *Chlamydomonas nivalis* by exposure to UV (Duval *et al.*, 2000) was accounted by stimulation of UV to the biochemical pathways related to proline metabolism.

The main amino acids in Antarctic microalgae changed in response to UV exposure, alanine, asparagine and glutamate increased after UV-B irradiation. The same results recorded in *U. lactuca* and *P. capillacea*, where alanine and glutamic acid increased after UV-B irradiation (Tables 4 & 6). The marked increase of alanine after exposure of *U. lactuca* and *P. capillacea* to UV-B might due to an enhancement of the alanine aminotransferase activity (Döhler *et al.*, 1997). Alanine decreased in *S. hornschurchii*, a result, which occurred at *Phaeocystis pouchetii* by its exposure to UV-B radiation, which was discussed by the damaging effect on the uptake of inorganic nitrogen and nitrogen metabolism (Döhler, 1992). Meanwhile, results found with UV sources regarding glutamine

and glutamate indicate a different influence on the glutamine synthetase/ glutamate synthase (GS/GOGAT) system (Döhler *et al.*, 1997).

Aspartic acid was reduced in all tested diatoms, a drastic reduction in glutamic acid could be observed in *L. annulata* samples (Döhler, 1984) which was discussed in relation to the impact of UV-B upon carbon and nitrogen metabolism. These data were in agreement with our results, where aspartic acid decreased in *U. lactuca* & *S. hornschurchii*. Döhler (1997) recorded that glutamine, serine and glycine decreased in antarctic microalgae by exposure to UV-B radiation. This was true for *S. hornschurchii*, where glutamic and glycine decreased due to UV-B irradiation (Table 5). The ¹⁵N-incorporation into the amino acids was reduced as a result of UV-B exposure of phytoplankton and ice algae. Results are discussed with reference to an inhibitory effect on the enzymes of both carbon and nitrogen metabolism as well as adaptation strategies (Döhler, 1997).

UV-B radiation is readily absorbed by nucleic acid and protein chromatophores, and their participation in plant responses to UV radiation has been documented (Buma *et al.*, 2003 and Jobson & Qiu, 2011). The involvement of these components in biological responses to UV radiation would indicate that protein synthesis and enzyme activities could be affected if biological systems were exposed to UV-B radiation (Garrard and Brandle, 1975). In addition, ultraviolet light can dimerize thymidine bases and cause lesions in DNA (Drake, 1970).

In the present study, the exposure of *U. lactuca* and *S. hornschurchii* to UV-B radiation caused the increase of protein content compared to the control as shown in Table 7. Exposure of *Scenedesmus quadricauda* to UV-A and UV-C caused increase in soluble proteins (Kovacik *et al.*, 2010) and *Dunaliella viridis* was found by Jiménez *et al.* (2004) to have the ability to adapt to a variety of environmental stresses including nitrogen starvation, osmotic or thermal shocks and UV irradiation by formation of proteins (50, 45 and 43 KDa) and increase its contents. Tominaga *et al.* (2010) found the accumulation of heat shock protein 70 (HSP70) in the alga *Ulva pertusa* by exposure to high temperature & suggested that this protein play a particularly important roles in adaptation to the stress conditions. UV-B exposure of higher plants leaves induces the synthesis of special polypeptids like stress-proteins (Santos *et al.*, 2004).

Kovacik *et al.* (2010) exposed axenic cultures of *Scenedesmus quadricauda* to UV-A (366 nm) and UV-C (254 nm) light over 1 h. Both wavelengths stimulated increase in soluble proteins. Primary photosynthetic carboxylating enzymes and soluble proteins in leaves of C3 and C4 crop plants were greatly affected by UV-B radiation (Vu *et al.*, 1982). Evidences suggest that polyamine accumulation may serve as indicator of UV radiation stress (Kramer *et al.*, 1991).

On the other hand, the protein content of *P. capillacea* decreased due to UV-B irradiation Table 7. This result was in agreement with those of some authors such as Noaman (2007) who noticed that the total proteins of *Synechococcus leopoliensis* exposed to UV decreased by increasing the exposure time. UV exposure for 24h caused the reduction of the protein content of *Dunaliella bardawil* (Salguero *et al.* 2005) and Bischof *et al.* (2000) noticed that UV

radiation resulted in loss of protein of some marine macroalgae. The same results obtained also by Bischof *et al.* (2000) who recorded that exposure to UV resulted in loss of total protein only in the deepwater species *Laminaria solidungula* and *Phycodrys rubens*. The different sensitivities to UV exposure of the species tested reflect their zonation pattern in the field.

Damage and degradation of protein by UV is proved in algae (Xue *et al.*, 2005). Kumar *et al.* (2003) proved the inhibition of nitrogenase enzyme by UV which may be the cause for inhibition of protein synthesis in *P. capillacea* or the damage may be due to the ability of protein to absorb UV which was proved by Ziska and Teramura (1992). Chaturvedi and Shyam (2000) proved that the degradation of protein of *Chlamydomonas reinhardtii* by its exposure to UV-B and Prasad *et al.* (1998) showed the inhibition of contents of protein by exposure *Chlorella vulgaris* to UV-B stress.

Formation of reactive oxygen species (ROS) in response to environmental stresses such as UV-B radiation is a common feature in plants (Rao *et al.*, 1996). In general, oxidative stress results from the disruption of cellular homeostasis of ROS production from the excitation of O₂ to form singlet (O_{1/2}) and the transfer of 1, 2 or 3 electrons to O₂ to form superoxide (O₂⁻), hydrogen peroxide (H₂O₂) and the hydroxyle radical (HO⁻), respectively (Halliwell and Gutteridge, 1989). The generation of ROS leads to oxidative destruction of the cell components through oxidative damage of membrane lipids, nucleic acid and protein (Imlay & Linn, 1998). To counteract the toxicity of ROS, defense systems that scavenge cellular ROS have been developed in plants to cope with oxidative stress via the non-enzymatic and enzymatic systems (Noctor and Foyer, 1998 and Asada, 1999). Antioxidants including water-soluble ascorbate (AsA) and water-insoluble α -tocopherol and carotenoids have been considered to be the nonenzymatic agents for scavenging ROS (Noctor & Foyer, 1998, Smirnoff & Wheeler, 2000 and Munné-Bosch & Alegre, 2002). In the enzymatic ROS-scavenging pathways, superoxide dismutase (SOD) converts O₂⁻ to H₂O₂ and then ascorbate peroxidase (APX) and glutathione reductase (GR) in the ascorbate-glutathione cycle (AGC) are responsible for H₂O₂ removal (Asada, 1999). Catalase (CAT) (Willekens *et al.*, 1997) and peroxidase (POX) (Asada and Takahashi, 1987) are also involved in H₂O₂ removal.

In the three irradiated species, *U. lactuca*, *S. hornschurchii* & *P. capillacea*, the activity of superoxide dismutase (Table 9), a prominent biomarker of defense against oxidative stress (Bowler *et al.*, 1992) increases with UV-B irradiation as a direct consequence. Fortunately antioxidant systems in plant and algae can scavenge ROS, including the antioxidant molecules such as carotenoids, ascorbate and reduced glutathione and antioxidant enzymes such as SOD, CAT, APX as well as several other enzymes involved in the ascorbate-glutathione cycle, which is considered to be an efficient ROS detoxifying system in chloroplasts (Jordan, 1996, Niyogi, 1999 and Foyer *et al.*, 1994). This was true for our results of APX (Table 8) and CAT (Table 10) in the three irradiated species, where the two enzymes of the antioxidant systems obviously increased due to UV-B radiation. It was demonstrated by Jansen *et al.*, (2001) that peroxidases are able to contribute

to the protection of PSII in plants from UV radiation stress (as a result of their oxygen radical scavenging activity through removing H₂O₂).

The over-production of ROS and the induction of oxidative stress by UV-B radiation have been observed in microalgae as *Chlorella vulgaris* (Malanga *et al.*, 1997), cyanobacteria (He *et al.*, 2002) and diatom (Rijstenbil, 2002). An increase in the activities of ROS scavenging enzymes was observed in algae exposed to oxidative stress (Rijstenbil, 2002). It is known that antioxidant defense mechanism against ROS is pivotal for algal survival under stressful conditions, higher antioxidant contents and antioxidant enzyme activities are associated with higher stress tolerance in algae (Collén & Davison, 1999 a, b). UV-B also increased APX and GR activities in *Pterocladia capillacea* but decreased them in *Gelidium amansii*. UV-B also increased SOD and CAT activities but to a higher degree in *G. amansii*. So, *G. amansii* suffered greater oxidative stress from UV-B radiation. *P. capillacea* can effectively reduce UV-B sensitivity by increasing sunscreen ability and antioxidant defense capacity (Lee and Shiu, 2009).

Additionally, the imbalance between light phase and Calvin cycle probably due to the decreased activity of ribulose-1,5-biphosphate carboxylase/ oxygenase (Rubisco) by UV irradiation (Bischof, 2000) promoted the formation of superoxide radical at the level of ferredoxin at photosystem I (PS I). The direct effect of UV-B on respiration pathway might contribute to the increased ROS formation. It is well-known that the overproduction of ROS in living organisms including photoautotrophs under stress conditions is potentially toxic which may attack biomolecules such as lipid, protein, DNA and some small molecules and results in oxidative damage, even the death of the organisms (Halliwell and Gutteridge, 1989).

SOD activity was increased in the marine macroalga *U. fasciata* by UV radiation (followed by a decrease at higher UV doses), which also increased the activities of CAT, POX, APX and GR. The induction of antioxidant enzyme activities for detoxifying reactive oxygen species (Shiu & Lee, 2005), which serves as the defense system against oxidative stress occurring in *U. fasciata* upon exposure to UV radiation. The excretion of H₂O₂ as well as the availability of antioxidants and the activation of SOD, CAT, guaiacol POX and reactive oxygen scavenging enzymes in the ascorbate-glutathione cycle serve as the defence system against oxidative stress occurring in *U. fasciata* upon exposure to UV-B. UV-B disrupts the balance between the production and removal of H₂O₂ and subsequently accumulated H₂O₂ initiates the signaling responses leading to the induction of enzymatic antioxidant defense systems to overcome ROS production in *U. fasciata* (Shiu and Lee, 2005).

Despite the fact that a large number of publications, especially during the last 20 years, is devoted to UV research in different algal systems, these mostly neglected to study possible influences on cell ultrastructure or the use of different transmission/scanning electron microscopy (TEM/SEM) methods to address structural changes in cellular components. Holzinger & Lütz (2006) postulated that UV-B effects on ultrastructure of algal cells can be found in one article in the book edited by Rozema *et al.* (1997) and a single communication by Lütz *et al.* (1997). The latter group used freshwater green algae also in another study on

ultrastructure changes and physiological adaptations under different stimulated UV regimes (Meindle and Lütz, 1996). More studies, presented as single communications, report on ultrastructure and UV-effects in marine diatoms (Buma *et al.*, 1996), Haptophyta (Buma *et al.*, 2000), marine red algae (Poppe *et al.*, 2002, 2003 and Holzinger *et al.*, 2004) and marine green algae (Holzinger *et al.*, 2006).

Exposure of the three species to extra doses of UV radiation caused the cells to show dissipation of the chloroplasts and irregularity in shapes (Plates 2, 4 & 6). *U. lactuca* showed disrupted chloroplast structure with severe damage in the thylakoid membranes when the alga irradiated for five days, 60 minutes daily. The same observation was also noticed, with different degrees, in chloroplasts of both *S. hornschurchii* and *P. capillacea*.

Holzinger *et al.* (2004) studied the effect of UV radiation on the ultrastructure of two red algae *Palmaria palmate* and *Odonthalia dentate*. Their TEM results demonstrated that the photosynthetic apparatus was severely influenced by UV, because thylakoid membranes appeared wrinkled, lumen dilatations occurred, and the outer membranes were altered. This dissipation of chloroplast was in full agreement with our results.

Poppe *et al.* (2002) reported destruction in chloroplasts, by exposure of the alga *Palmaria decipiens* to UV for 8 hours. UV irradiation of *Palmaria palmata* for 6 hours caused damage in the outer chloroplast envelope and lumen dilatation (Holzinger *et al.*, 2004). Meanwhile irradiation for 24 hours caused severe damage with irregular lumen of the thylakoids. As the same time severe damage in the thylakoid membranes occurred after 24 hours exposure of the red alga *Odonthalia dentata* to UV irradiation (Holzinger *et al.*, 2004).

Under UV-B stress, the thylakoid membrane of *Spirulina platensis* becomes distorted (Gupta *et al.*, 2008). It must be mentioned that the thylakoid membrane is the site for both photosynthesis and respiration (Gantt, 1994). Structural disturbance to membranes is likely to result in a reduced photosynthetic activity, e. g. due to dilation of the thylakoid membranes and rupture of the chloroplast double membrane (Strid *et al.*, 1994).

The chloroplast envelope and the thylakoids membranes of the red macroalga, *Phycodrys austrogeogica* were damaged and the phycobilisomes were detached from the thylakoids after 12 hours UV irradiance (Poppe *et al.*, 2003). The same study showed that in the red alga *Palmaria decipiens*, UV irradiation for 4 hours lead to changes in ultrastructure of chloroplasts, with dilated thylakoids as compared to the control cell, while 6 and 8 hours irradiation caused disrupted thylakoids and the formation of inside-out translucent thylakoids vesicles, which also shown in irradiated *S. hornschurchii* as shown in plate 7B. Changes to the ultrastructure of chloroplast due to UV treatment and a vesiculation of the thylakoids was observed. Drastic changes in the arrangement of thylakoids membranes were found and a large number of small plasma vesicles accumulated at the plasma membrane as a consequence of UV irradiation (Holzinger *et al.*, 2004).

The effect of ultraviolet (UV) radiation on the ultrastructure of four red algae, the endemic Antarctic *Palmaria decipiens* and *Phycodrys austrogeogica*,

the Arctic-cold temperate *Palmaria palmate* and the cosmopolitan *Bangia atropurpurea* was studied. All four species showed a formation of 'insideout' vesicles from the chloroplast thylakoids upon exposure to artificial UV-radiation. In *P. decipiens*, most vesicles were developed after 8 h and in *P. palmate* after 48 h of UV exposure. In *B. atropurpurea*, vesiculation of thylakoids was observed after 72 h of UV irradiation. In *Ph. Austrogeorgica*, the chloroplast envelope and thylakoid membranes were damaged and the phycobilisomes became detached from the thylakoids after 12 h of UV exposure (Poppe *et al.*, 2003).

Buma *et al.* (1996) proved that the irradiated diatoms *Cyclotella sp.*, *Nitzschia colsterium* and *Thalassiosira nordenskioldii* by UV radiation showed that vacuolization had taken place, the initial large vacuole was found to be fragmented into small vacuoles and the nuclear envelope as well as membranes appeared to be unaffected by UV irradiation in the three diatoms. The same results were showed by Holzinger *et al.* (2004), where no changes in nuclei or Golgi bodies in *Palmaria decipiens* treated by UV irradiation, and the nuclear membrane of *Palmaria palmate* appeared normal, which demonstrates that membranes are not generally destroyed but selectively targeted and altered by UV treatment.

The above discussion was in consistence with the ultrastructure of the investigated species, where, as shown in plates (2, 4 & 6) the nucleus of *U. lactuca* was damaged and showed wrinkled nuclear envelope due to UV-B irradiation. On the other hand, the nucleus in *S. hornschurchii* and *P. capillacea* is nearly unaffected. At the same time some vacuoles were appeared in the three investigated species due to UV-B irradiation, where *U. lactuca* contained a small vacuole, meanwhile *S. hornschurchii* and *P. capillacea* showed appearance of some vacuols. UV-B irradiation also affected the cell wall of the three species, since *U. lactuca* showed lamellated cell wall, while *S. hornschurchii* and *P. capillacea* showed irregularity of their walls.

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