

Supplementary Figures

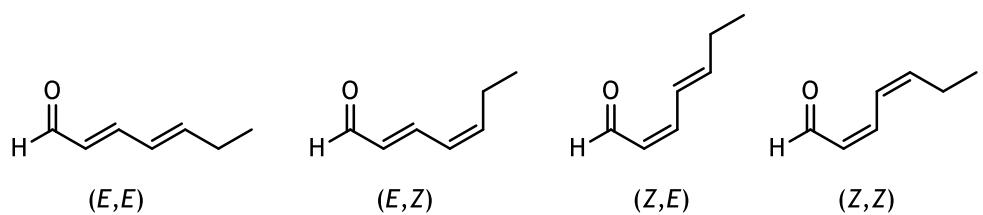


Figure S1: Molecular structure of isomers Type I using heptadienal as an example.

Supplementary tables

Table S1: Phytoplankton functional groups used for FLOW CAM quantification of large size phytoplankton (10-250 µm) cell abundance and biovolume. Grouping name for analysis is detailed.

Category	Size range 09 to 20 µm	Size range 20 to 100 µm	Size range 100 to 250 µm
Others	Spherical cells	Spherical cells	
	Elliptical cells	Elliptical cells	Elliptical cells
	Discoid cells	Discoid cells	
	Cylindrical cells		
	Filamentous chains	Filamentous chains	Filamentous chains
Coccolithophorids		Coccolithophores	
Silicoflagellates	Silicoflagellates	Silicoflagellates	Silicoflagellates
Dinoflagellates		<i>Protoperidinium</i>	<i>Protoperidinium</i>
		<i>Ceratium</i>	<i>Ceratium</i>
		<i>Prorocentrum</i>	<i>Prorocentrum</i>
		<i>Gymnodiniales</i>	<i>Gymnodiniales</i>
		<i>Peridiniales</i>	
		Unidentified dinoflagellates	Unidentified dinoflagellates
Tintinids		Extended-lorica tintinnids	Extended-lorica tintinnids
		Globular-lorica tintinnids	Globular-lorica tintinnids
Diatoms		Pennate diatoms	Pennate diatoms
		Needle-like diatoms	Needle-like diatoms
		<i>Rhizosolenia & Proboscia</i>	Proboscia
		Wide-chain diatoms	<i>Pleurosigma</i>
		<i>Guinardia striata</i>	
		<i>Blavyanus flaccida</i>	
		<i>Leptocylindrus</i>	
		Filamentous diatoms	
		Slightly-connected diatoms	
		<i>Chaetoceros</i>	<i>Chaetoceros</i>
		<i>Skeletonema</i>	<i>Skeletonema</i>
		Unidentified single centric diatoms	Unidentified single centric diatoms
		Unidentified short-chain diatoms	
		Unidentified long-chain diatoms	
		<i>Thalassiosira</i>	
		<i>Asterionellopsis</i>	<i>Asterionellopsis</i>
		<i>Thalassionema</i>	<i>Thalassionema</i>
Phaeocystis		<i>Phaeocystis</i>	

Table S2: Spearman rank correlation coefficients performed on particulate (pPUA) and dissolved PUA (dPUA) data and environmental variables. d: dissolved; p: particulate; C7: 2E,4E/Z-heptadienal. C8: 2E,4E/Z-octadienal; C10: 2E,4E/Z-decadienal; T: Temperature ($^{\circ}\text{C}$); S: Salinity; PAR: Photosynthetic Active Radiation ($\text{mmol quanta m}^{-2} \text{s}^{-1}$); F: Fluorescence (f.u.); ϵ : coefficient of dissipation of turbulent kinetic energy ($\text{m}^2 \text{s}^{-3}$); Nitrate, phosphate and Silicate: μM . O_2 : mg L $^{-1}$.

* significant at level 0.05: ** significant at level 0.01.

Station depth	PUA	T	S	PAR	F	NO_3^-	PO_4^{3-}	SiO_4	ϵ	Oxygen
Jet 5 m	dPUA	0.206	-0.181	0.034	0.180	-0.140	-0.058	-0.186	0.598**	0.198
	dC7	0.259	-0.169	0.08	0.136	-0.195	-0.028	-0.162	0.333	0.227
	dC8	0.182	-0.204	-0.022	0.200	-0.137	-0.074	-0.206	0.660**	0.193
	dC10	0.169	-0.119	0.072	0.133	-0.094	-0.039	0.068	0.522**	0.15
	pPUA	0.55	0.67	-0.458	0.748	-0.317	0.189	-0.335	0.099	-0.26
	pC7	0.77	0.78	-0.411	0.755	-0.267	-0.061	-0.57	0.13	-0.25
	pC8	0.13	0.32	-0.505	0.631	-0.580	0.577	0.131	0.03	-0.28
	pC10	0.41	0.68	-0.260	0.575	-0.164	0.103	-0.371	0.059	-0.08
	dPUA	-0.041	0.056	0.287	-0.104	0.116	0.251	-0.425	-0.133	-0.0023
	dC7	0.26	-0.17	0.08	0.156	-0.060	0.218	-0.133	-0.085	0.188
Coast 5 m	dC8	0.182	-0.204	-0.022	-0.128	-0.103	0.278	-0.041	-0.126	0.05
	dC10	-0.141	0.17	-0.12	0.073	0.346	0.083	0.105	-0.152	-0.34
	pPUA	-0.955	-0.219	0.662	-0.060	0.400	-0.322	0.568	-0.94	0.318
	pC7	-0.917	-0.173	0.519	-0.129	0.569	-0.480	0.463	-0.89	0.177
	pC8	-0.917	-0.226	0.854	0.016	0.102	-0.061	0.655	-0.921	0.484
	pC10	-0.751	-0.486	0.821	0.410	-0.406	0.476	0.848	-0.791	0.827
	dPUA	-0.148	0.125	0.0126	-0.05	0.132	0.044	0.113	0.359	-0.196
	dC7	-0.17	0.128	0.03	-0.04	0.141	0.076	0.101	0.454	-0.184
	dC8	-0.07	0.08	0.004	-0.06	0.081	-0.028	0.097	0.236	-0.173
	dC10	-0.278	0.217	-0.01	-0.05	0.223	0.168	0.160	0.283	-0.239
Gyre 5 m	pPUA	-0.012	0.01	0.902	0.227	-0.424	-0.455	-0.219	-0.109	-0.029
	pC7	-0.852	-0.796	0.287	-0.266	-0.497	-0.730	-0.158	0.75	0.816
	pC8	-0.194	-0.171	0.897	0.089	-0.516	-0.529	-0.202	0.046	0.14
	pC10	0.234	0.245	0.892	0.338	-0.310	-0.290	-0.199	-0.245	-0.267
	pPUA tot	0.55	0.67	-0.458	0.748	-0.317	0.189	-0.335	0.099	-0.26
	pC7	0.77	0.78	-0.411	0.755	-0.144	-0.061	-0.570	0.13	-0.25
	pC8	0.13	0.32	-0.505	0.631	-0.580	0.577	0.131	0.03	-0.28
	pC10	0.41	0.68	-0.26	0.575	-0.164	0.103	-0.371	0.059	-0.08
	dPUA	0.206	-0.181	0.034	0.18	-0.140	-0.058	-0.186	0.598	0.198
	dC7	0.259	-0.169	0.08	0.136	-0.195	-0.028	-0.162	0.333	0.227
Jet DCM	dC8	0.182	-0.204	-0.022	0.2	-0.137	-0.074	-0.206	0.660	0.193
	dC10	0.169	-0.119	0.072	0.133	-0.094	-0.039	-0.131	0.522	0.150
	pPUA	-0.955	-0.219	0.662	-0.06	0.400	-0.322	0.568	-0.940	0.318
	pC7	-0.917	-0.173	0.519	-0.129	0.569	-0.480	0.463	-0.890	0.177
	pC8	-0.917	-0.226	0.854	0.016	0.102	-0.061	0.655	-0.921	0.484
	pC10	-0.751	-0.486	0.821	0.41	-0.426	0.476	0.848	-0.791	0.827
	dPUA tot	-0.041	0.056	0.287	-0.104	0.116	0.251	-0.425	-0.133	-0.002
	HD	0.123	-0.131	0.128	0.156	-0.060	0.218	-0.133	-0.080	0.188
	OD	0.039	-0.003	0.544	-0.128	0.069	0.278	-0.041	-0.120	0.050
	DD	-0.36	0.362	-0.103	-0.339	0.346	0.083	0.105	-0.152	-0.340
Gyre DCM	pPUA tot	-0.012	0.01	0.902	0.227	-0.424	-0.455	-0.219	-0.109	-0.029
	HD	-0.852	-0.796	0.287	-0.266	-0.497	-0.730	-0.158	0.750	0.816
	OD	-0.194	-0.171	0.897	0.089	-0.516	-0.529	-0.202	0.046	0.140
	DD	0.234	0.245	0.892	0.338	-0.310	-0.290	-0.199	-0.245	-0.267
	dPUA tot	-0.148	0.125	0.0126	-0.05	0.132	0.044	0.113	0.359	-0.196
	HD	-0.17	0.128	0.03	-0.04	0.141	0.076	0.101	0.454	-0.184
	OD	-0.07	0.08	0.004	-0.06	0.081	-0.028	0.097	0.236	-0.173
	DD	-0.278	0.217	-0.01	-0.05	0.223	0.168	0.160	0.283	-0.239

Table S3: Spearman rank correlation coefficients performed on pPUA data and biological variables I (pigments) at the DCM of the different sites. High correlation are highlighted in grey. dPUA: dissolved PUA; pPUA: particulate PUA; dC7: 2E,4E/Z-heptadienal. dC8: 2E,4E/Z-octadienal; dC10: 2E,4E/Z-decadienal. Perid.: peridinin; Fx: Fucoxanthin; Viol.: violaxanthin; Prasin.; prasinoxanthin; Diadin.; diadinoxanthin; Allox.; alloxanthine; But-Fx: 19'-butanoyloxyfucoxanthin. Hex-Fx: 19'-hexanoyloxyfucoxanthin. Significant values are highlighted in bold. “–” denotes undetected concentrations.

Station	pPUA	Chla	Chlc ₁	Chlc ₂	Chlc ₃	Perid.	But-Fx	Fx	Violax	Prasinox	Hex-Fx	Diadin.	Allox	Diatox.	Luteine
Jet	pPUA	0.91	-0.11	0.42	0.079	-0.256	0.003	0.918	0.441	0.977	0.964	0.901	0.842	-0.381	0.62
	pC7	0.946	-0.60	0.63	0.031	-0.023	-0.511	0.954	0.183	0.785	0.704	0.973	0.991	-0.414	0.502
	pC8	0.31	0.709	-0.07	0.217	-0.394	0.766	0.265	0.463	0.631	0.724	0.224	0.096	0.011	0.311
	pC10	0.51	-0.29	-0.21	-0.64	-0.432	-0.089	0.792	0.777	0.775	0.758	0.683	0.586	-0.896	0.991
Coast	pPUA	0.937	0.911	-0.982	0.783	0.03	0.587	0.851	0.65	0.533	0.984	0.803	0.99	0.908	
	pC7	0.988	0.809	-0.999	0.643	-0.171	0.413	0.728	0.485	0.352	0.929	0.668	0.986	0.806	
	pC8	0.769	0.996	-0.865	0.945	0.361	0.823	0.977	0.866	0.784	0.986	0.956	0.929	0.995	
	pC10	0.475	0.956	-0.616	0.998	0.683	0.975	0.986	0.99	0.959	0.854	0.996	0.725	0.958	
Gyre	pPUA	-0.214		-		-0.374	-0.456		-		-0.227	-0.351	-	-0.141	-
	pC7	0.494		-		-0.193	0.178		-		0.188	0.481	-	0.31	-
	pC8	-0.047		-		-0.33	-0.34		-		-0.107	-0.191	-	-0.004	-
	pC10	-0.385		-		-0.36	-0.554		-		-0.314	-0.527	-	-0.256	-

Table S4: Spearman rank correlation coefficients performed on pPUA data and biological variables II (biovolume of taxonomical categories of large size phytoplankton) at the DCM of the different sites. High correlation is highlighted in grey. Significance level * $p<0.05$; ** $p<0.01$. dPUA: pPUA: particulate PUA; dC7: 2E,4E/Z-heptadienal. dC8: 2E,4E/Z-octadienal; dC10: 2E,4E/Z-decadienal. Significant values are highlighted with grey area.

	Diatoms	Cocolitophorids	Silicoflagellates	Dinoflagellates	Others
Jet	pPUA	0.730	-0.425	0.881*	0.043
	pC7	0.831*	-0.326	0.897*	0.057
	pC8	0.463	-0.536	0.622	0.117
	pC10	0.568	-0.292	0.945**	-0.215
Coast	pPUA	-0.086	-0.157	0.244	-0.041
	pC7	-0.174	-0.208	0.339	-0.189
	pC8	0.021	-0.106	0.026	0.171
	pC10	0.446	0.286	-0.029	0.645*
Gyre	pPUA	-0.559	-0.43	0.092	-0.483
	pC7	-0.193	-0.429	0.106	-0.687
	pC8	-0.591	-0.524	0.109	-0.611
	pC10	-0.542	-0.33	0.068	-0.318
Total	pPUA	0.338	-0.079	0.569*	-0.103
	pC7	0.503*	0.085	0.744*	-0.073
	pC8	0.333	-0.103	0.566*	-0.058
	pC10	0.578*	0.0109	0.746*	-0.065