

Microalgae as sustainable bio-factories of healthy lipids: evaluating fatty acid content and antioxidant activity

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Table S1. Lipid content of *Chlorella vulgaris*, *Chlorococcum amblystomatis*, *Scenedesmus obliquus*, *Tetraselmis chui*, *Phaeodactylum tricornutum*, *Spirulina* sp. and *Nannochloropsis oceanica*. Kruskal–Wallis test followed by Dunn’s post-hoc comparisons).

Microalgae	<i>Chlorophyta</i>				<i>Bacillariophyta</i>	<i>Cyanobacteria</i>	<i>Ochrophyta</i>
	<i>Chlorella vulgaris</i>	<i>Chlorococcum amblystomatis</i>	<i>Scenedesmus obliquus</i>	<i>Tetraselmis chui</i>	<i>Phaeodactylum tricornutum</i>	<i>Spirulina</i> sp.	<i>Nannochloropsis oceanica</i>
Lipid content (%)	8.8±0.7 ^{a,b}	16.6±0.9 ^{a,c}	11.1±1.1	6.5±0.4 ^{c,d,e}	12.6±1.2 ^e	10.7±1.0 ^f	20.9±3.4^{b,d,f}

Table S2. – Evaluation of the antioxidant activity for the extracts of different microalgae. Concentration of lipid extract ($\mu\text{g.mL}^{-1}$) providing 50% inhibition of the ABTS^{•+} radical and 20% inhibition of the DPPH[•] radical, and their respective Trolox equivalents (TE; $\mu\text{mol.g}^{-1}$). The values are displayed as the mean (n = 3) ± standard deviation. Matching letters (a–f) indicate statistically significant differences between microalgae species, i.e. the same letter represents significant differences ($q < 0.05$, Kruskal–Wallis test followed by Dunn’s post-hoc comparisons).

		<i>Chlorophyta</i>				<i>Bacillariophyta</i>	<i>Cyanobacteria</i>	<i>Ochrophyta</i>
		<i>C. vulgaris</i>	<i>C. amblystomatis</i>	<i>S. obliquus</i>	<i>T. chui</i>	<i>P. tricornutum</i>	<i>Spirulina</i> sp.	<i>N. oceanica</i>
ABTS^{•+}	IC50 ($\mu\text{g.mL}^{-1}$)	51.1 ±3.7	52.6±4.6	29.4±1.2 ^b	40.9±4.7	57.3±4.5 ^b	38.7±1.6 ^c	101.9±1.7 ^{a,c}
	TE ($\mu\text{mol.g}^{-1}$)	368.4 ±27.0	358.8±32.4	637.5±27.4 ^a	432.9±51.3	306.6±23.5	452.2±9.6	184.0±3.2 ^a
DPPH[•]	IC20 ($\mu\text{g.mL}^{-1}$)	50.5 ±12.3 ^{d,e}	58.4±10.7 ^f	89.1±6.6	225.7±6.9 ^{d,f}	75.4±4.6	96.9±9.7	175.6±8.7 ^e
	TE ($\mu\text{mol.g}^{-1}$)	191.8 ±40.0 ^{d,e}	143.0±25.1 ^f	114.5±8.9	45.0±1.4 ^{d,f}	122.3±7.2	84.3±9.1	52.5±2.7 ^e

A. Hierarchical clustering

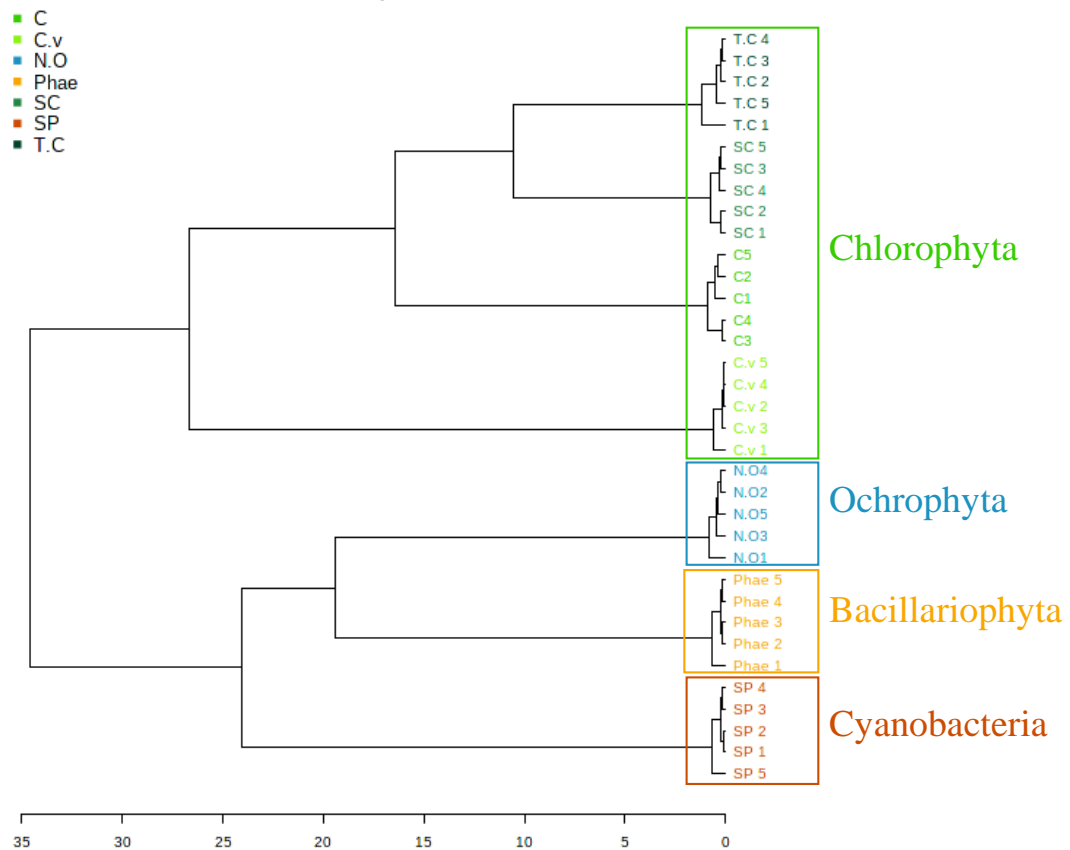


Figure S1. (A) Hierarchical cluster analysis using relative abundance after Glog normalization of all fatty acids identified in *Chlorella vulgaris*, *Chlorococcum amblystomatis*, *Scenedesmus obliquus*, *Tetraselmis chui*, *Phaeodactylum tricornutum*, *Spirulina* sp. and *Nannochloropsis oceanica*. The green, blue, yellow, and orange boxes show microalgae from the phylum Chlorophyta, Ochrophyta, Bacillariophyta and Cyanobacteria, respectively. Abbreviations: C, *Chlorococcum amblystomatis*; C.v, *Chlorella vulgaris*; SC, *Scenedesmus obliquus*; T.C, *Tetraselmis chui*; Phae, *Phaeodactylum tricornutum*; SP, *Spirulina* sp.; and N.O, *Nannochloropsis oceanica*.

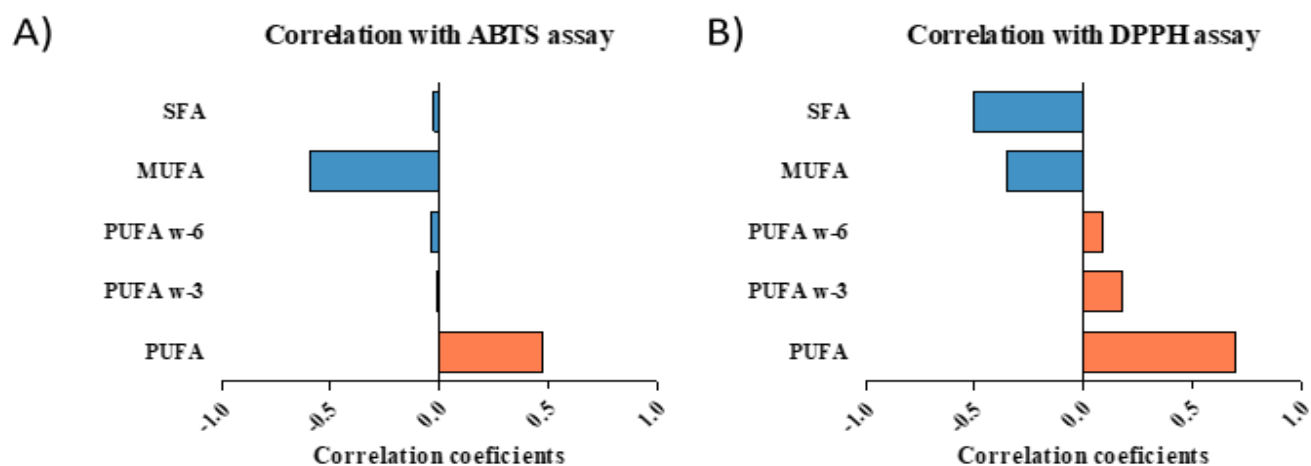


Figure S2. Correlation coefficients analysis using the sum of relative abundances from saturated fatty acids (SFA), mono-unsaturated FA (MUFA), polyunsaturated FA (PUFA), omega-3 PUFA and omega-6 PUFA, and antioxidant activity. **(A)** Correlation coefficients with ABTS assay. **(B)** Correlation coefficients with DPPH assay.