



Article Fatty Acid Changes in Nearshore Phytoplankton under Anthropogenic Impact as a Biodiversity Risk Factor for the World's Deepest Lake Baikal

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Copyright: © 2022 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (https:// creativecommons.org/licenses/by/ 4.0/). Abstract: In this study, we present results on fatty acid analysis of phytoplankton of Lake Baikal, the world's deepest lake, which differs from other lakes by its oceanic features. Since we used a large-mesh net, the net sample phytoplankton were primarily represented by the large elongated diatom Synedra acus. subsp. radians (Kützing) Skabichevskij. The similar algae composition of net samples of spring season phytoplankton collected at different sites of the lake allows us to compare results of the fatty acid analysis of these samples. The phytoplankton diversity of the sedimentation samples was contrary represented by 32 algae species. There are clear changes in the fatty acid composition of net phytoplankton exposed to anthropogenic impacts of varying intensity. The content of polyunsaturated fatty acids in phytoplankton collected from central stations (pelagic stations at a distance of \sim 10–30 km from the shoreline) without anthropogenic impact was higher by up to 15% than phytoplankton collected from nearshore stations (littoral stations at a distance of $\sim 0.01-0.05$ km from the shoreline) and offshore stations (pelagic stations at a distance of ~ 3 km from the shoreline). The interlaboratory precision of fatty acid determination of phytoplankton is estimated as $\leq 10\%$. We found high content of the lipid peroxidation marker (80–340 µg g⁻¹ of dry weight) in phytoplankton from nearshore and offshore stations with intensive anthropogenic impact. In phytoplankton from central stations, we did not find any lipid peroxidation. Determination of unsaturated fatty acids, coupled with analysis of fatty acid peroxidation products, can be used to evaluate the level of anthropogenic impact in terms of ecological health and biodiversity conservation.

Keywords: Lake Baikal phytoplankton; *Synedra acus*; fatty acids; peroxidation; nearshore; pelagic; PUFA; anionic surfactants; oxidative stress; linear alkylbenzene sulphonates

1. Introduction

Lake Baikal is the world's deepest (maximum depth of 1642 m) [1,2] and oldest (20–25 million years) lake, containing 23,615.39 km³ of fresh water with a mineralization of 96–98 mg L⁻¹. Lake Baikal, a rift lake, is located in the southern part of Eastern Siberia (Russia) and comprises three basins: northern, central, and southern [3,4].

There are both littoral and pelagic zones in Lake Baikal [3,5,6]. The lake is 636 km long, with a maximum width of 79 km. Its littoral zone reaches 80–120 m of depth, including the nearshore zone, which extends to a 25 m depth [7,8]. The tectonic terrace on the western flank rises over the lake surface at heights of about 300–1500 m and continues steeply underwater to 1200–1400 m. It then turns into the valley bottom, which slants toward the east coast [9]. As demonstrated in a schematic bathymetric map (Figure 1), the platform is fairly shallow near the shore but drops off steeply, especially on the western flank. This



platform step forms a boundary between the littoral and pelagic zones. The surface area of Lake Baikal is $32,822 \text{ km}^2$, of which the littoral zone occupies only ~3.4%.

Figure 1. Bathymetric map of Lake Baikal. Sampling sites are marked.

Nevertheless, the nearshore zone of Lake Baikal is the most productive, containing more than 98% of all bottom dwellers, such as Porifera, Polychaeta, Oligochaeta, and other worms, along with Sagina, Isopoda, Trichoptera, Chironomidae, Mollusca, and fishes. Their biomass in the southern basin was 620 kg per hectare at depths of \leq 4 m and 100–150 kg per hectare at depths of 20–70 m [10,11].

Phytoplankton are the primary producers of organic matter in Lake Baikal, inhabiting the photic zone at depths of ~60–120 m. Intensive photosynthesis takes place in this zone. Phytoplankton also inhabit the epipelagic zone, a zone with depths of ~200–300 m and maximum vertical mixing of water [10]. Microalgae live in the deep-water pelagic zone

down to 750 m of depth [12] without cell division due to mixotrophic growth [13] and the rest stage of growth during adverse conditions [14]. Some microalgae and their spores drop to the bottom water layer during the disadvantageous season, floating to the surface later in the spring period. After blooming, phytoplankton become food for protozoa, zooplankton, benthic organisms, and bacteria, helping maintain biodiversity. Spring primary production of Lake Baikal phytoplankton reaches ~160 tons per hectare (i.e., ~80% of the annual biomass) [10]. Phytoplankton are highly sensitive to environmental changes, and their health benefits the well-being of the entire lake ecosystem [15–18].

The high productivity and biodiversity in the nearshore zone of Lake Baikal are due to unique oceanic or marine features of climate, hydrological regime, and bathymetry. In oceans (and seas), there are places where the ocean floor deepens closer to shore. In coastal areas of oceans, the interaction of winds and currents generates vertical movement of cold, nutrient-rich deepwater towards the surface (upwelling). As in oceans, upwelling is a typical phenomenon for Lake Baikal [19,20]. Causes of vertical water movement include asymmetric depression and the presence of high-pressure thermal and cold-water springs through bottom cracks. Another cause is climate features and winds formed due to the great length of Lake Baikal from south to north, which leads to a baric gradient between the northern and southern basins. Winds also cause movement in the water column down to the bottom [21,22]. The most intensive upwelling processes and nutrient enrichment take place near the western and northern coasts of Lake Baikal [21–24]. These several peculiarities coupled with the runoff of ten largest tributaries of the lake (the total water discharge of ~1522 m³ s⁻¹) lead to an almost unlimited concentration of nutrients in the coastal zone, which is sufficient for the autotrophic organisms providing the bioproductivity of the lake [24–26].

For Baikal phytoplankton, the bulk of primary production comprises lipids [27]. The fatty acid (FA) composition of phytoplankton depends on metabolism, abiotic factors [28–30], and biotic factors. Lake Baikal phytoplankton produce esterified and free fatty acids [29], including essential polyunsaturated fatty acids (PUFA) [31,32]. FAs migrate through food webs and have an impact on the functioning and energetic potential of all organisms [33,34]. Changes in phytoplankton FA composition can elucidate the ecosystem stress caused by anthropogenic impact [35–37].

For live cells, which are the structural units of whole organisms, the cell wall is the first target of negative anthropogenic impacts. Phospholipids are dominant constituents of the cell wall. There are two known routes for lipid destruction in the cell. The first is the α -, β -, and ω -oxidation of lipids by enzymes with the formation of numerous vital compounds, including acetyl coenzyme A (acetyl-CoA) with its further oxidation to adenosine triphosphate (ATP) and energy generation. The second is lipid peroxidation. One final product of peroxidation is malondialdehyde (MDA), which can exist in the cell in free (f-MDA) and covalently-bonded (b-MDA) forms [38–40]. MDA reacts with lipids, thiol groups, and amino groups of nucleic acids of proteins, as well as with lipoproteins and DNA, resulting in the destruction of these biological molecules [34–36]. MDA can also react with DNA to form mutagenic adducts [41,42]. Lipid peroxidation products, including MDA, can be analyzed in their reaction with thiobarbituric acid, at which point they are called thiobarbituric reactive substances (TBARS).

In large aquatic ecosystems, the nearshore zone is much more exposed to negative anthropogenic impact than the pelagic zone [15] and can result in stress. The narrow nearshore zone of Lake Baikal, with its unique biodiversity, warrants concern. Changes in littoral phytoplankton from 2000 to 2015 were observed [43]. For example, the biomass of atypical small unicellular green algae and atypical filamentous green algae increased significantly. On the other hand, psychrophilic pelagic phytoplankton did not change significantly from 1964 to 2011. Diatoms comprised about 50–90% of biomass [44], and changes in the structure of benthic communities [45] and the death of Lake Baikal sponges in the littoral zone were also observed [46,47].

Understanding the dynamics and functioning of nearshore communities is necessary to decrease the anthropogenic impact on the littoral zone. This study aims to reveal stress in Baikal phytoplankton through the analysis of FAs and their peroxidation products as a response of the environment to anthropogenic impacts of different intensities.

2. Experiment

2.1. Sample Collection

For Lake Baikal, a delay of seasonal onsets in the coastal zone is common compared to the nearby continental zone [24]. Phytoplankton samples were collected in the spring seasons of 2014, 2016, 2018, and 2020, from 28 May to 20 June, using the vertical Juday-type net with 100 μ m mesh size. Samples were collected from a depth of 20 m at the coastal and pelagic stations located in the three Lake Baikal basins (Figure 1).

All samples were collected from the stations of three types. These types differ by their distances from shore. The differences are the following.

1. Pelagic phytoplankton were collected from central stations located within the central deep-water areas of the three basins. Of all stations, these were the farthest from shore. The distance from the shoreline was ~10–30 km. The central stations no. 11, 13, 15, 17, and 18 were considered to lack anthropogenic impact.

2. Pelagic phytoplankton were collected from offshore pelagic stations located ~3 km from the shoreline. The offshore stations no. 10, 12, 14, and 16 were considered to have a low anthropogenic impact.

3. Coastal phytoplankton were collected from the stations located about 0.01–0.05 km from the shoreline. These are considered nearshore stations. Some were located near cities (no. 0, 2, 3, and 9) and settlements (no. 1, 5, and 8). The nearshore stations no. 4, 5, and 6 were located in areas with intensive recreation. Nearshore stations no. 4 and 7 were located far from any settlements.

For the spring season, the phytoplankton *Bacillariophyta* and *Dinophyta* microalgae are typical. For example, the mean cell length of the dominant *Synedra acus subsp. radians* (Kützing) Skabichevskij (this name is currently regarded as a synonym of *Fragilaria radians* (Kützing) D.M. Williams and Round) reaches 142–288 μ m, and the length of some cells reaches 500 μ m. These algae dominated from 2014 to 2020. *Aulacoseira baikalensis* (Wislouch) Simonsen, a diatom endemic to Lake Baikal, forms filaments up to 160–1000 μ m long, but this species was not dominant from 2014 to 2020. We used a 100 μ m mesh net to obtain *S. acus* as net samples, excluding other small algae because different algae species have various responses to stress factors. This allows us to compare the results of chemical analysis of phytoplankton samples obtained in different parts of the lake. Samples containing zooplankton organisms were filtered through a nylon net with 400 mesh size to separate them from phytoplankton. Phytoplankton net samples were immediately filtered onto glass fiber filters (47 mm), wrapped in aluminum foil, and stored at -20 °C before chemical analysis.

The separate samples of phytoplankton were collected to analyze Baikal phytoplankton diversity by the classical sedimentation method [6]. Water from the surface down to a depth of 1 m was sampled using the bathometer. A water sample of 1 L volume was poured into a plastic bottle, which was kept in dark at 20 °C. For the estimation of algal biomass, samples were fixed with Lugol's solution. The sedimentation process lasted 10 days. Supernatant was removed by drop method using a siphon. It was shown previously that there were no algae cells in each drop of the removed supernatant. Samples then underwent concentration to 0.5–1 mL. Cells in concentrated sedimentation samples were subsequently identified and counted using light microscopy (Axiovert 200, Carl Zeiss, Jena, Germany) at 400× magnification. According to microscopic analysis, zooplankton content was negligible in all net samples (<0.5%). The number of cells was converted into phytoplankton biomass, taking into account the biovolume measurements of individual cells. Mean cell dimensions were used to estimate cell biovolumes for different species. The separate water samples were collected at coastal (n = 53) and pelagic (n = 4) sites in the three basins to analyze anionic surfactant concentrations. Surface water to a depth of 1 m was sampled in autumn 2019, from 13 to 22 September, and in the spring season 2020, from 28 May to 10 June, using the bathometer.

2.2. Lipid Extraction and Derivatization of FA to Fatty Acid Methyl Esters (FAMEs)

Wet sample weights ranged from 0.15–0.20 g. The humidity of the samples was estimated gravimetrically, with values reaching ~90%. To extract lipids from replicate samples, 1.2 mL of Folch solution (chloroform–methanol, 2:1, by volume) was added to each sample. Samples then underwent extraction for 5 min using a 50 Hz ultrasonic bath, after which 0.35 mL of distilled water was added to the extracts. The mixtures were vigorously shaken for 0.5 min and centrifuged at 13,000 rpm for 3 min. The chloroform extracts were put into 10 mL glass vials and the solvent evaporated using an argon stream [31].

Fatty acid methyl esters (FAME) were derived through the acidic esterification of fatty acids. First, 4.5 mL of 2% H₂SO₄ solution in methanol was added to the dry fraction. The bottles were covered with aluminum foil caps and kept for 1.5 h at 55 °C. Next, 0.8 mL of *n*-hexane was added to the FAME solutions and the samples were cooled to room temperature. FAMEs were extracted twice with *n*-hexane (3 mL × 2 min). The extracts were first dried with anhydrous Na₂SO₄ and then concentrated to 1 mL volume using argon stream. An internal standard was added to the extract.

2.3. FAME Analysis by Gas Chromatography with Flame-Ionization (GC-FID) and Mass-Spectrometric Detection (GC-MS)

For qualitative and quantitative analyses, the internal standard (1 mg mL⁻¹ of dodecyl ether solution in *n*-hexane) and external standard solutions of FAs "FAME Mix C4–C24" and "methyl cis-4,7,10,13,16,19-docosahexaenoic acid" solution (both Supelco, USA) were used. Qualitative analysis of FAMEs by GC-MS was carried out using the Agilent 6890B GC System coupled to the 7000C GC/MS Triple Quad mass spectrometer (Agilent, USA). The injector temperature was 290 °C; the injection volume was 2 μ L in splitless mode; the quadrupole temperature was 150 °C; the ion source temperature was 230 °C; the ionization energy was 70 eV. Chromatographic peaks were detected in the *m/z* 40–500 range. FAMEs were identified based on retention times and their mass spectra using the NIST Mass Spectral Search Program for the NIST/EPA/NIH Mass Spectral Library software (V. 2.2).

Quantitative analysis of FAMEs was carried out using a Shimadzu GC-2010-Plus gas chromatograph (Shimadzu, Japan). Injection of 2 μ L of sample into the column was carried out in the splitless regime using an Optima-17MS GC column (30 m × 0.25 mm, SGE Macherey-Nagel, Germany). The injector temperature was 310 °C; the injection volume was 2 μ L in splitless mode; the detector temperature was 310 °C; the hydrogen velocity was 40 mL/min; the air velocity was 400 mL/min; the carrier-gas velocity (helium) was 30 mL/min; the linear velocity was 28.5 cm/s. Chromatography was carried out by heating the column first from 80 °C to 120 °C at the rate of 10 °C/min and then to 260 °C at the rate of 2.5 °C/min.

2.4. Fatty Acid Peroxidation Estimate by Spectrophotometric Determination of Thiobarbituric acid Reactive Substances (TBARS)

Thiobarbituric acid reactive substances (TBARS) can form pink TBA₂-TBARS adducts with thiobarbituric acid (TBA) with an absorption maximum at 532 nm, which is a true indicator of lipid peroxidation [38,48,49]. In our study, we analyzed TBARS using the modified technique. Wet sample weights ranged from 0.15–0.20 g at ~85–90% humidity. Samples were placed in plastic, 2-mL Eppendorf taste-tubes, and 1 mL of 10% trichloroacetic acid (TCA) solution was added. Samples were sonicated for 5 min in a 50 Hz ultrasonic bath and centrifuged at 13,000 rpm for 3 min. The supernatant was put into 5 mL glass vials with plastic screw caps and mixed with 2 mL of 0.5% TBA solution in 10% TCA. The mixture was kept for 0.5 h at 95 °C, cooled, and centrifuged. Absorbance was measured using a Cintra-20 spectrophotometer (GBC, Australia) and standart quartz quvettes (path

length 1 cm). The TBARS extinction coefficient was 155 mM⁻¹ L. Baseline absorption was measured at 532 nm using the solution obtained by mixing 1 mL of 10% TCA and 2 mL of 0.5% TBA solutions. The TBARS absorption was measured at 532 nm. The limit of quantification (LOQ) of 1.21 μ mol mL⁻¹ extract and the limit of determination (LOD) of 0.51 μ mol mL⁻¹ via spectrophotometry were determined by Rakita et al. [50].

2.5. Statistical Data

The laboratory precision of FAME determinations was estimated by calculating the relative standard deviations (RSD) of two replicate results from the average value using the degree of freedom number f = 22. For selected subgroups, the homogeneity of variations was checked using Fisher and Cochran tests [51]. The repeatability of the technique for total FAME content was characterized by V = 6%. We also used three replicates for the in vitro microalgae growing experiment.

For the determination of similarities and differences between natural samples and laboratory cultures in terms of FA concentrations and profiles, the principal component analysis of multidimensional statistics was used [52]. Results were visualized using the *Statistica* software.

To determine similarities between the fatty acid spectra of samples, cluster analysis was carried out using the *Statistica* software. To estimate distances between stations, Ward's method based on multifactorial variance analysis was used. The matrix of distances was calculated using the Euclidean method.

2.6. Cultivation of the S. acus

S. acus diatoms were grown in 100 cm³ Erlenmeyer flasks at +6 °C for 50 days with the chemically defined medium (DM medium) containing Ca(NO₃)₂ × 4H₂O (20 mg L⁻¹), KH₂PO₄ (12.4 mg L⁻¹), MgSO₄ × 7H₂O (25 mg L⁻¹), NaHCO₃ (16 mg L⁻¹), Na₂EDTA (23 mg L⁻¹), H₃BO₃ (2.48 mg L⁻¹), MnCl × 4H₂O (1.4 mg L⁻¹), (NH₄)₆Mo₇O₂₄ × 4H₂O (1.0 mg L⁻¹), cyanocobalamin (0.04 mg L⁻¹), thiamine hydrochloride (0.04 mg L⁻¹), biotin (0.04 mg L⁻¹), Na₂SiO₃ × 5H₂O (42.6 mg L⁻¹), and FeCl₃ (1.6 mg L⁻¹). The 12/12 h day/night light cycle was provided by luminescent bulbs with a photon flux density of ~200 μ E m⁻² s⁻¹. Biomass sampling was carried out over 12 h, 24 h, and 14 days.

2.7. Anionic Surfactant Analysis by Spectrophotometric Determination

Anionic surfactants were analyzed as a complex with methylene blue dye. Linear alkylbenzene sulphonate (LAS) mixture was used as an external standard. The calibration equation y = 3.0423x + 0.0041 ($R^2 = 0.9933$) was obtained in the range of concentrations from 2 to 100 µg L⁻¹. LAS were extracted from the 0.05 L water samples with trichloromethane (5 mL × 2 min). All water samples passed through cellulose acetate filters (45 µm, VLADiS-ART, Russia) using the filter-apparatus (Duran Group, Germany) to separate suspended matter on filters. Preliminary extraction of surfactants absorbed on suspended matter such as phytoplankton was carried out with ethanol (1 mL × 2 min). Both soluble and absorbed surfactants were involved in the extraction. Absorbance was measured using a Cintra-20 spectrophotometer with quartz quvettes (path length of 1 cm). The methylene blue and anionic surfactant complex has a maximum absorbance at 651.5 nm.

3. Results

3.1. Baikal Phytoplankton Composition

Microscopy data revealed that the *S. acus* diatom dominated most samples of littoral and pelagic Baikal phytoplankton during the spring seasons of 2014, 2016, 2018, and 2020. The composition of Baikal phytoplankton (2020) is presented in Table 1.

	Sample Stations in South Basin of the Lake									
Algae Taxa	Western Flank Nearshore Sampling Site No. 19			Pelagic Sampling Site No. 20					Eastern Flank Nearshore Sampling Site No. 14	
				Sampling Depth						
	1	m	1 r	n	50	m	400	0 m	1	m
		Nur	nber of cel	ls N, 10 ³	cells L ⁻	¹ and cel	l bioma	ss B, mg i	m^{-3}	
	N	В	Ν	В	Ν	В	Ν	В	Ν	В
		Cl	ass Cyanoj	phyta						
Anabaena sp.	_	-	-	-	3.4	0.34	-	-	-	_
<i>Oscillatoria</i> sp.	42	5.9	-	-	-	-	-	-	-	-
<i>Lyngbya</i> sp.	-	-	-	-	_	-	-	-	-	-
Phormidium sp.	_	-	-	-	_	_	-	-	-	-
Total	42	5.9	-	-	3.4	0.3	-	-	-	_
		Clá	ass Chryso	phyta						
Dinobryon cylindricum O.E. Imhof	0.2	0.4	0.3	0.6	1.4	2.7	2.5	4.8	0.4	0.20
Cystes of Chrysophyta	0.6	0.3	0.6	0.3	1.4	0.7	6.5	3.3	-	_
<i>Dinobryon sociale</i> (Ehrenberg) Ehrenberg	-	-	-	-	-	-	-	-	-	-
Mallomonas vannigera Asmund	_	-	-	-	-	-	-	-	98	4.9
Chrysochromulina parva Lackey	58	2.9	42	2.1	8.4	0.4	0.5	0.03	-	-
Total	59	3.6	43	3.0	11	3.8	10	8.0	98	5.1
		Cla	ass Crypto	phyta						
Rhodomonas pusilla (H. Bachmann) Javornicky	65	16	26	6.3	17	4.0	2.2	0.5	145	35
Cryptomonas gracilis Skuja	0.4	0.8	0.6	1.2	1.1	2.2	0.5	1.1	0.6	1.2
Cryptomonas ovata Ehrenberg	_	-	-	-	_	_	-	-	_	-
Total	65	16	27	7.5	18	6.3	2.7	1.6	146	36
		С	lass Dinop	hyta						
<i>Gyrodinium helveticum</i> (Penard) Y. Takano and T. Horiguchi	0.4	10	0.9	23	2.3	58	1.1	28	0.4	11
<i>Gymnodinium baicalense</i> N.L. Antipova	0.2	6	0.15	4.5	_	-	_	_	0.20	6.0
<i>Peridinium baicalense</i> Kisselev and V. Zvetkov	_	-	-	_	_	_	-	-	-	-
Glenodinium sp.	0.4	0.6	0.9	1.3	1.1	1.7	2.7	4.1	1.4	2.0
<i>Ceratium hirundinella</i> (O.F.Müller) Dujardin	_	-	-	_	_	_	-	-	-	-
Total	1.0	17	1.9	28	3.4	59	3.8	32	2.0	19
		Clas	ss Bacillari	ophyta						
Aulacoseira baikalensis (Wislouch) Simonsen	0.4	6	_	_	0.3	4.5	0.5	8.1	3.2	48
<i>Aulacoseira islandica</i> (O.Müller) Simonsen + spore	0.8	4	0.6	3	2	10	3.8	19	34	170

Table 1. Comprehensive composition of the current spring season phytoplankton in LakeBaikal (2020).

Chlamidomonas sp.

Total phytoplankton biomass

Total

	Sample Stations in South Basin of the Lake									
Algae Taxa	Western Flank Nearshore Sampling Site No. 19		Pelagic Sampling Site No. 20						Eastern Flank Nearshore Sampling Site No. 14	
				S	amplin	g Depth				
<i>Synedra acus subsp. radians</i> (Kützing) Skabichevskij	131	249	210	399	140	266	65	124	280	532
<i>Synedra ulna var. danica</i> (Kützing) Van Heurck	0.2	0.6	_	-	-	-	0.5	1.7	-	_
Synedra ulna (Nitzsch) Ehrenberg	0.1	0.22	-	_	_	_	-	_	_	_
<i>Nitzschia graciliformis</i> Lange-Bertalot and Simonsen	_	_	0.3	0.08	_	_	_	-	-	_
<i>Cyclotella minuta</i> (Skvortsov) Antipova	0.2	0.4	_	_	0.3	0.6	0.3	0.6	0.20	0.4
<i>Cyclotella baicalensis</i> Skvortsov and K.I.Meyer	0.2	4.0	0.3	5.8	0.3	6	0.3	6.0	0.20	4.0
Ellerbeckia teres (Brun) R.M.Crawford	-	-	-	-	_	-	-	-	-	-
<i>Stephanodiscus meyeri</i> Genkal and Popovskaya	0.2	0.2	0.6	0.6	0.6	0.6	0.8	0.8	-	_
Asterionella formosa Hassall	-	-	-	-	_	-	-	-	-	-
Total	133	264	212	408	144	288	71	160	318	754
		Cla	ass Chloro	ophyta						
Monoraphidium arcuatum (Korshikov)'Hindák	15	1.8	8.6	1.0	13	1.6	0.8	0.10	6.2	0.7
Monoraphidium contortum (Thuret) Komárková-Legnerová	_	-	_	_	0.8	0.11	_	-	-	_
Koliella longiseta (Vischer) Hindák	0.3	0.08	0.6	0.15	0.3	0.08	0.8	0.20	0.7	0.18
Elakatothrix genevensis (Reverdin) Hindák	-	-	0.14	0.02	-	-	-	-	-	-
<i>Scenedesmus acuminatus</i> (Lagerheim) Chodat	-	-	-	-	-	-	-	-	-	_
Scenedesmus sp.	-	-	_	-	-	-	-	-	-	-
<i>Closterium moniliferum</i> Ehrenberg ex Ralfs	_	_	_	_	-	-	_	-	-	_
Pediastrum boryanum (Turpin) Meneghini	-	-	_	_	-	-	_	-	-	_

_

1.9

309

9.3

293

16

316

Table 1. Cont.

In all sedimentation samples of littoral and pelagic phytoplankton (2014, 2016–2021), *S. acus* subsp. *radians* dominated. Six algal classes represented the phytoplankton composition of sedimentation samples: Cyanophyta, Chrysophyta, Cryptophyta, Dinophyta, Bacillariophyta, and Chlorophyta (Table 2). As shown, Bacillariophyta were dominant during the spring season, with biomass values from 79% in deep water to 93% in surface water.

_

14.1

194

1.2

448

_

1.7

359

0.3

1.9

89

0.15

0.4

202

_

6.9

571

0.9

815

All net samples of littoral and pelagic phytoplankton contained *S. acus* as a dominant species in 2014 and 2016. *S. acus* contribution to total biomass was ~95%, excluding small-celled phytoplankton. The large-volume algae (Figure 2) *A. baicalensis, Aulacoseira islandica* (O. Müller) Simonsen *and Ceratium hirundinella* (O.F. Müller) Dujardin, long-size *Synedra ulna* (Nitzsch) Ehrenberg and *Synedra ulna var. danica* (Kützing) Van Heurck, small-volume *Cyclotella baicalensis* Skvortsov and K.I. Meyer and *Cyclotella minuta* (Skvortsov) Antipova, both of which may form long chains, and *Dinobryon cylindricum* O.E. Imhof, all contributed less than 5% to the total biomass of net phytoplankton.

	Sample Stations in the Southern Basin of Lake Baikal												
Class of Algae	W Flank S	Vestern Nearshore Station		Eastern Flank Nearshore Station									
				Sampling Depth									
	1 m 1 m			m	50 m			0 m	1	m			
	Number of cells N (%) and cell biomass B (%)												
	Ν	В	Ν	В	Ν	В	Ν	В	Ν	В			
Cyanophyta	13	1.9	-	-	1.8	0.09	-	-	-	-			
Chrysophyta	19	1.2	15	0.7	5.8	1.1	11	4.0	17.2	0.6			
Cryptophyta	21	5.3	9.2	1.7	9.3	1.7	3.1	0.8	26	4.4			
Dinophyta	0.3	5.4	0.7	6.3	1.8	16	4.3	16	0.4	2.3			
Bacillariophy	/ta 42	86	72	91	74	80	80	79	56	93			
Chlorophyta	4.9	0.6	3.2	0.3	7.3	0.5	2.1	0.22	1.2	0.11			

Table 2. Contribution of Bacillariophyta to total phytoplankton biomass during spring season (2020).



Figure 2. Some large-volume and long-sized algae in phytoplankton of Lake Baikal during the spring season 2020. Sample site no. 20.

The compositions of net samples and sedimentation samples were different. All sedimentation samples of littoral and pelagic phytoplankton contained *S. acus* as a dominant species in 2014 and 2016. *S. acus* contribution to the total biomass was up to 95% (Table 3).

Sampling Stations	Northern Latitude (N)	Eastern Longitude (E)	B (%) of S. acus	Other Predominant Bacillariophyta
0	51°31′391″	104°11′398″	91	A. baicalensis, C. baicalensis, A. islandica
1	51°43′147″	103°42′964″	95	A. baicalensis, C. baicalensis, A. islandica
2	51°31′726″	104°08′086″	91	A. baicalensis, C. baicalensis, A. islandica
3	51°31′726″	104°08′086″	91	A. baicalensis, C. baicalensis, A. islandica
4	52°47′332″	106°36′445″	85	A. islandica
5	53°12′316″	107°19′587″	87	A. islandica
6	53°13′201″	107°22′290″	87	A. islandica
7	54°32′285″	108°39′814″	89	A. baicalensis, A. islandica
8	55°21′428″	109°12′713″	88	A. baicalensis, A. islandica
9.1	55°35′385″	109°20′720″	88	A. baicalensis, A. islandica
9.1	55°34′021″	109°13′677″	88	A. baicalensis, A. islandica
19	51°49′094″	104°54′776″	95	A. baicalensis, C. baicalensis, A. islandica

Table 3. Contribution of *S. acus* to total coastal phytoplankton biomass during spring season (2014).

3.2. Statistical Analysis

The comparative statistical analysis of data on FA content in the samples of the axenic culture of *S. acus* and natural plankton dominated by *S. acus* using the principal component method is shown in Figure 3. This method takes into account a high number of individual FAs in every sample. It allows us to visualize data through the conversion of the multidimensional space to a two-dimensional one. As the principal components, we selected FAs of different saturations and chain lengths (1–22 carbon atoms). These FAs are C16:0, C16:1(ω-7), C16:2(ω-4), C16:3(ω-4), C18:0, C18:1(ω-9), C18:3(ω-3), C18:4(ω-3), C20:4(*w*-6), C20:5(*w*-3), and C22:6(*w*-3). Of total FA weight, 70–80% belongs to these FAs. Percentages of the aforementioned FAs in laboratory culture biomass at stationary and exponential phases of growth, as well as in the coastal (no. 1 to 9) and pelagic phytoplankton samples (no. 10 to 17), were chosen as variables in the data array. Three separate projections to the factor plane were made for laboratory culture (Figure 3, top), coastal phytoplankton (Figure 4, bottom left), and pelagic phytoplankton variables (Figure 3, bottom right). The significant validity (\geq 90%) of presented results was marked for all three cases. Based on analysis of the FA composition of the laboratory culture determined by this statistical method, differences are clear for diatoms at stationary and exponential growth phases (Figure 3, top).

The vectors of most samples of the coastal Baikal phytoplankton lie in the upper left-half plane, except for samples no. 4, 6, and 8. This figure demonstrates the significant difference between the nearshore samples no. 4, 6, and 8 from the cities and other coastal samples.



Figure 3. Analysis of Baikal phytoplankton and *S. acus* axenic laboratory culture by the principal component method.

Considering that samples were collected at the spring blooming peak, we can explain with relative confidence the positioning of vectors for the samples of coastal phytoplankton (Figure 3, bottom left), which is similar to that of vectors for laboratory culture samples at the stage of exponential growth. Concerning pelagic Baikal phytoplankton, the maximum closeness between FA compositions of the samples from central pelagic stations no. 11, 13, 15, and 17 that were maximally remote from the shore is noteworthy. FA composition of these central samples differed from offshore pelagic ones.

Cluster analysis of the samples is presented (Figure 4). There are two main clusters of the samples, which do not cross. Laboratory culture samples at the stage of exponential growth are clustered together with pelagic samples and the bulk of coastal samples. This can be explained by the fact that during the sampling of natural phytoplankton, all stations were at the stage of intensive cell division. Moreover, it is also obvious that the pelagic plankton samples and the bulk of the coastal plankton samples form the two main clusters and show clear differences in FA composition of coastal and pelagic plankton (Figure 4).



Figure 4. Dendrogram of the FA composition of the Baikal coastal and pelagic phytoplankton and the FA composition of the *S. acus* laboratory culture at stationary and exponential phases of growth.

The difference between the phytoplankton from the central pelagic stations and the offshore pelagic stations is possibly related to the distance from the shore. Stations no. 11, 13, 15, and 17 were located in the central area of the lake in contrast to stations no. 10, 12, 14, and 16, which were located at a distance of 3 km from the shore (Figures 1 and 4). Some pelagic and coastal samples clustered together with laboratory culture samples at the stationary phase (Figure 4). This suggests that at these stations, natural phytoplankton were the same as at the stationary phase, which contradicts the microscopy data. Therefore, it is necessary to carefully compare the results of laboratory experiments and those of natural phytoplankton. Data analysis using the principal component method and the cluster analysis method is related to the so-called exploring analysis. This allows us to find dependencies in the analysis of large data arrays, but their interpretation is beyond the scope of mathematical statistics [53].

3.3. FA Composition and FA Peroxidation Product Analysis

In our study, we analyzed the FA composition and TBARS concentration in natural phytoplankton as well as in laboratory culture of *S. acus*. Total FA content in the pelagic and nearshore phytoplankton samples (2014 and 2016) varied from 20–60 mg g⁻¹ of dry weight (d.w.). Myristic C14:0, palmitic C16:0, and oleic C18:0 acids were dominant saturated FAs (SFAs) of ~30–40%. Palmitoleic acid C16:1(n-7) was the dominant MUFA (~20–30%). Hexatrienoic C16:3(n-6,9,12) and essential eicosapentaenoic C20:5(n-5,8,11,14,17) acids were dominant PUFAs (~30–40%). For all samples, high contents of C16:1 and C16:0 FAs (26–48%), as well as of C20:5 ω -3 FA (13–23%), were observed (Table 4).

	FA Composition, %								
		Phytoplankton	Labora	Laboratory S. acus Culture					
FAs	Nearshore Stations Far from Settlements	Central Pelagic Stations	Nearshore Stations Near Settlements	Offshore Pelagic Stations	4 h	24 h	2 Weeks		
C16:1 w9	30.4	21.7	19.2	12.2	12.0	13.3	27.0		
C16:0	17.3	11.4	17.4	13.6	37.2	36.1	21.0		
C16:3 w4	5.5	14.2	4.5	10.8	4.8	6.0	8.4		
C16:2 w4	3.6	4.3	2.9	4.7	2.2	2.9	3.9		
C18:1 w9	2.3	1.9	2.3	0.7	5.6	5.4	1.7		
C18:0	5.1	1.5	5.1	12.8	19	16	5.5		
C18:4 w3	2.4	0.2	2.4	0.8	1.65	2.11	2.4		
C18:3 w3	1.7	0.3	1.7	0.5	1.55	1.33	1.04		
C20:4 w6	0.3	0.3	0.1	0.3	0.08	0.05	0.02		
C20:5 w3	12.9	23.0	13.3	14.6	6.6	8.9	13.0		
C22:6 w3	3.8	1.7	2.0	3.7	0.59	1.02	1.27		
PUFAs	30.3	44.1	27.6	35.4	10.1	14.7	31.9		
PUFAs + MUFAs *	65	72	53	50	37	43	68		

Table 4. FA content averages of natural phytoplankton and laboratory culture of S. acus.

* PUFA—polyunsaturated fatty acids; MUFA—monounsaturated fatty acids.

The most similar FA compositions were observed for nearshore phytoplankton collected far from settlements and *S. acus* cultivated at 6 °C in poor medium during the two-week period (Table 4). The maximum content of polyunsaturated FAs (PUFAs) and monounsaturated FAs (MUFAs), including essential C20:5 ω 3 eicosapentaenoic acid, was recorded in pelagic phytoplankton from the central stations (Table 3).

The formation of malondialdehyde (MDA) as an FA peroxidation product characterizes pathological processes in a living cell. MDA can appear in water organisms under the impact of anionic surfactants, such as linear alkylbenzene sulphonates (LASs) [41,42,54], heavy metals [55], polycyclic aromatic hydrocarbons (PAH) [56], and antibiotics [57]. According to a previous investigation [58], anionic surfactants found in the Baikal nearshore water ranged from 4.8 μ g L⁻¹ to 20 μ g L⁻¹ [58]. Low concentrations of LASs (5–20 μ g L⁻¹) can cause toxic effects on aquatic microorganisms [59–61].

Based on these data, we analyzed for the first time the lipid peroxidation marker in Lake Baikal phytoplankton and laboratory cultures of *S. acus*. In phytoplankton collected from central pelagic stations no. 11, 13, 15, 17, and 18 and at nearshore stations no. 4 and 8, TBARS were not detected (Table 5). In phytoplankton of nearshore station no. 6, TBARS were represented a minimal concentration (14 μ g g⁻¹ of d.w.). TBARS concentration in phytoplankton of other nearshore stations near the Kultuk settlement, as well as near towns of Baikalsk and Severobaikalsk, ranged from 80–340 μ g g⁻¹ of d.w. TBARS were either not found or represented a minimal concentration in samples with high PUFA and MUFA percentages of ~70%. A TBARS concentration of \geq 80 μ g g⁻¹ was identified in samples with an average PUFA and MUFA content of 53 μ g g⁻¹ of d.w. In the *S. acus* laboratory culture, both at exponential and stationary stages of growth, TBARS were not detected (Table 5).

Samples	s Location	Lake Zone	Total FA, mg g ⁻¹	PUFA and MUFA Percentage	TBARS, μg g ⁻¹ of d.w.
11	Marituy River—Solzan River (2016)	Pelagic	15.8 ± 1.4	72	Not found
17	Baikalskoe settlement—Frolikha River (2016)	Pelagic	17.0 ± 1.5	72	Not found
18	Elokhin Cape—Davsha settlement	Pelagic	24.2 ± 2.2	70	Not found
1	Kultuk settlement (2016)	Nearshore	24.0 ± 2.1	56	80
1	Kultuk settlement (2018)	Nearshore	20.7 ± 1.8	56	100
2	Baikalsk town (2016)	Nearshore	27.1 ± 2.4	58	280
3	Baikalsk town (2016)	Nearshore	27.1 ± 2.4	58	340
0	Baikalsk pulp and paper mill (2016)	Nearshore	26.2 ± 2.3	60	160
0	Baikalsk pulp and paper mill (2018)	Nearshore	21.3 ± 1.9	54	164
4	Aya Bay (2016)	Nearshore	28.1 ± 2.5	66	Not found
6	Shamanka Bay (2016)	Nearshore	28.7 ± 2.6	60	14
7	Elokhin Cape (2016)	Nearshore	18.1 ± 1.6	55	180
8	Baikalskoe settlement (2016)	Nearshore	49 ± 4	68	Not found
9.1	Tyya River mouth (2016)	Nearshore	26.7 ± 2.4	46	100
9.1	Tyya River mouth (2018)	Nearshore	15.6 ± 1.4	58	11
9.2	Senogda Bay (2016)	Nearshore	14.0 ± 1.3	40	190
S. acus la	aboratory culture, +6 °C, DM medium, 12 l	n	4.2 ± 0.4	48	Not found
S. acus la	aboratory culture, +6 °C, DM medium, 24 l	n	35 ± 3	74	Not found
S. acus la	aboratory culture, +6 °C, DM medium, 50 o	days	No d	lata	Not found

Table 5. TBARS concentrations ($\mu g g^{-1}$ of d.w.) in some phytoplankton samples collected in 2016.

4. Discussion

4.1. The Present Composition of Baikal Phytoplankton during the Spring Season

Previously, it was thought that during the spring season the algal complex of pelagic phytoplankton in Lake Baikal consisted predominantly of large-volume diatoms [44], such as endemic *Aulacoseira baicalensis* (Wislouch) Simonsen, *A. islandica* (O.Müller) Simonsen, *Cyclotella minuta* (Skvortzov) Antipova, endemic *Cyclotella baicalensis* Skvortzov et Meyer; *S. acus* subsp. *radians* (Kützing) Skabitschevsky, and *Stephanodiscus meyeri* Genkal et Popovskaya. A similar phytoplankton composition is typical of pelagic and coastal zones [54–56]. Notably, for the past period (~30 years), the structure and functioning of shallow spring phytoplankton have changed. The periodicity of years with high productivity ($B \ge 1000 \text{ mg m}^{-3}$) and low productivity ($B \le 100 \text{ mg m}^{-3}$) of large-volume *A. baicalensis* has disappeared [62,63]. In the last ten years, the small-volume diatom *S. acus* subsp. *radians* has been maximally abundant during each spring season. Phytoplankton productivity in the current period from 2014 to 2021 was much lower than in previous years (1950–1970 and 1970–1995) when large-cell diatoms of the genus *Aulacoseira* predominated [12,63,64]. In recent years, *A. baicalensis* has been found as single cells without intensive blooming.

Our data show that in all littoral and pelagic net samples, phytoplankton (2016–2020) containing *S. acus*, prevailed. This allows us to compare the FA composition of *S. acus* fraction of coastal and pelagic phytoplankton exposed to anthropogenic impacts of varying intensity. The dominance of *S. acus* is needed to compare the results of FA analysis of the samples collected within different parts of the lake. This is due to *S. acus* prevalence and its sensitivity to pollution. We found that *S. acus* is a bioindicator of anionic surfactant pollution—it responds to the exposure of anionic surfactants of 5–20 μ g L⁻¹ concentration.

The response appears as oxidation stress due to a high content of PUFAs in cell membranes of diatoms. Diatom membranes consist of proteins and phospholipids (~50%/50% by mass) including PUFAs. For example, the PUFA content of *S. acus* cell membrane is ~30%. On the other hand, the cell membranes of Chrysophyta, Cryptophyta, Dinophyta, Cyanophyta, and Chlorophyta classes consist mostly of polymers. These polymers such as cellulose and hemicellulose make these algae more stress resistant.

Many lakes show vertical stratification of their water masses, at least for some extended periods. In Lake Baikal, we can observe inverse water stratification in March, with a surface water temperature of ~0.4–2 °C [64]. In the deep-water layer, the water temperature is ~3.6 °C. At the end of May or in the first two weeks of June, we observed spring homothermy of the water body, during which water temperature is ~3.5–3.7 °C through all layers, from the bottom to the surface [65]. At the end of August or in the first two weeks of September, we can observe a reverse water stratification, in which surface water temperature is ~3.5–3.7 °C, and thermocline depth is 25–50 m. At the end of November, an autumn homothermy can occur. In our study, all samples were collected during the spring season when water temperatures were near homothermy values [64].

In our investigation, water temperature ranged from 3.4–3.6 °C at almost all pelagic sampling sites and from 3.0–6.0 °C at almost all coastal sites. Nitrate, phosphate, and silica concentrations in Baikal water are constant throughout the year [10,19,66]. According to Popovskaya et al., there is no clear relationship between the concentration of nutrients and phytoplankton growth [44]. The seasonal by-turn blooming of the different species is mostly due to temperature changes. The temperature during the spring season increases irregularly throughout the entire water area of the lake due to its large length from south to north. According to Popovskaya et al. [44], there is no clear relationship between the intensity of phytoplankton growth to water temperatures of 0.8 °C to 5.0 °C. Hence, our study did not take into account water temperatures and nutrient concentrations that provide high productivity in coastal and pelagic zones [10,21,25,26] and which cannot cause stress or change in phytoplankton FA composition under spring season homothermy.

4.2. Phytoplankton FA Composition and FA Peroxidation

At all sampling sites, phytoplankton were at the exponential stage of growth, allowing us to exclude the dependence of FA composition on growth stages. Multivariate analysis of the results by the principal component method and the cluster analysis confirms that FA composition of pelagic and coastal phytoplankton differs.

A higher PUFA concentration (~40%) was determined in the samples of pelagic phytoplankton compared to most of the coastal samples (~26%). Additionally, a higher content (up to 10–15%) of MUFA of triacylglycerols was determined in the coastal samples. Higher concentrations of the sum of MUFA and PUFA (~65%) were detected in coastal samples no. 4, 6, and 8 compared to other coastal samples (~53%). We noted differences in PUFA concentrations between pelagic phytoplankton collected at the central stations and phytoplankton from the offshore pelagic stations. Trying to understand this difference, this study analyzed, for the first time, TBARS as a lipid peroxidation marker in Baikal phytoplankton and laboratory cultures of *S. acus*. Table 5 shows that TBARS were either not found or shows minimal concentrations in the coastal samples no. 4, 6, and 8 samples. Aya Bay (station no. 4), Shamanka Bay (station no. 6), and Ludar Cape (Station no. 8), where the Baikalskoe settlement is situated (population 600 people), are the areas with low anthropogenic load compared to other nearshore stations. Oxidative stress was not identified at these stations. As mentioned above, the principal component method and cluster analysis revealed the clustering of these samples separately from other coastal samples (Figures 3 and 4).

Coastal stations no. 1, 2, 3, 5, and 9 were situated near large cities and settlements, such as Kultuk (3715 citizens, according to 2020 data), Baikalsk (12,534), Khuzhir (1350), and Severobaikalsk (23,180). The high TBARS concentration in phytoplankton of these

stations ranged from 80–340 μ g g⁻¹ of dry weight (Table 5). High TBARS concentration near Elokhin Cape can be explained by cyclonic convectional gradient flow along the perimeter of the lake, as this flow can transport water from the Severobaikalsk town (station no. 9) along the west coast to Elokhin Cape [23,67].

Therefore, we suggest that this phenomenon of TBARS occurrence in phytoplankton results from peroxidation of cell membrane phospholipids (mainly PUFAs) due to anthropogenic impact in the coastal zone. We assume that this impact can be caused by anionic surfactants. These substances were found in Lake Baikal for the first time in 2019 in our previous investigation [58]. According to the mentioned investigation, anionic surfactant concentrations in the coastal water of Lake Baikal ranged from 4.8 μ g L⁻¹ (station no. 8) to 20 μ g L⁻¹ (station no. 9). A higher concentration of linear alkylbenzene sulphonates LAS (40 μ g L⁻¹) was found in the Tyya River, a tributary of Lake Baikal. The Tyya mouth is located close to the station no. 9 (~3 km). A dramatically high concentration of anionic surfactants (170 μ g L⁻¹) was registered in sewage water of Severobaikalsk (Republic of Buryatia) [57]. We also analyzed anionic surfactants in surface water of three basins (n = 57) using a modified spectrophotometry method with methylene blue and gas chromatography coupled with mass spectrometry. Surfactant concentrations ranged from 4.5 ± 0.6 to 29.2 ± 3.5 . μ g L $^{-1}$ in samples collected in autumn 2019, and from 2.00 ± 0.24 to $16.9 \pm 2.1 \,\mu g \, L^{-1}$ in samples collected in spring season 2020. Maximum concentrations $(\geq 20 \ \mu g \ L^{-1}, n = 3)$ were found in the recreation zones as well as in zones located not far from settlements.

There is no production plant at the coast of Lake Baikal dumping untreated sewage water into the lake. Nevertheless, the recreation and human impacts on the environment have increased significantly [68]. The dramatic effect relates to the absence of wastewater treatment plants on the Lake Baikal coast. Few treatment plants are of outdated design. As a result, eutrophication of local nearshore areas as well as the toxic effect of environmental pollutants of untreated sewage water can be observed. Anionic surfactants such as LASs are the most common aquatic pollutants. Low LAS concentrations (5–20 μ g L⁻¹) have toxic effects on aquatic microorganisms [59,60] and fishes, leading to morphological changes in gills and skin [61]. Given this, we can assume a negative impact of low surfactant concentrations on Baikal phytoplankton. This phenomenon can cause changes in the production of biologically active substances including fatty acids that puts the biodiversity of the world's deepest lake at risk. Baikal is one of the largest and deepest lake of the world natural heritage of the UNESCO. It differs from other lakes by its unique marine features. A lot of fine poetic words like these "Hoc mare [Yonder is the sea], magnum et spatiosum manibus [great and wide]; illic reptilia quorum non est numerus [Wherein are things creeping innumerable], animalia pusilla cum magnis [Both small and great beasts]" can be dedicated to great lakes. But nearshore living organisms of great water ecosystems need our efficient actions to decrease anthropogenic impact on littoral. Phytoplankton represent the first link of the water ecosystem. Their damage by chemicals may result in serious biodiversity changes, the onset of which can already be observed in the nearshore zone.

5. Conclusions

For the first time, we revealed a lipid peroxidation product (TBARS) in the Baikal nearshore phytoplankton collected near large cities and settlements (80–340 μ g g⁻¹). TBARS were not detected in the phytoplankton of stations that were maximally remote from the sources of anthropogenic impact. Changes in fatty acid composition, including a decrease in PUFA, were identified for phytoplankton exposed to anthropogenic impact. This suggests that the high level of anthropogenic impact on the coastal zone near cities and large settlements causes stress of phytoplankton.

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