A symbiosis of microorganisms with sponges is not only one of the World’s most ancient but also one of the most difficult, combining more than 100 species of organisms from various taxonomic groups into a single symbiotic community. Symbionts play an important role in the biotransformation of organic matter. Therefore, studies of adaptive features in an animal organism for the representatives of taxa at a low phylogenetic level are currently very relevant in terms of the ever-increasing anthropogenic pressure on stable ecosystems.

The anthropogenic impact has also changed the functioning of many ecosystems in Lake Baikal. In recent years, there has been a massive death of the community of endemic Baikal sponges (Timoshkin et al., 2019).

The cause of this phenomenon can be determined through a study of a sponge community under laboratory conditions. Previous studies allowed for analysis of the variability of the lipid, pigment and protein composition of the cells of the Baikal sponges (Latyshev et al., 1992; Glyzina and Glyzin, 2014; Itskovich et al., 2017; 2019; Glyzina et al., 2017). Thus, we identified taxonomic differences in the composition of fatty acids in the Baikal sponges and their symbionts and studied the influence of some factors (habitat and light intensity) on chlorophyll-porphyrins, lipids and fatty acids of sponges. We revealed that despite the variability of the fatty acid composition, the features typical of the Baikal sponges are preserved.

During this study, a symbiotic community based on the Baikal endemic sponge *Lubomirskia baicalensis* (Pallas) (Efremova, 2004) served as a study object, including microalgae, bacteria and unicellular organisms living in sponge cells (Itskovich et al., 2019). The sponge was collected from a depth of 10 m in the southern part of Lake Baikal and adapted to artificial habitat in aquarium installations at a temperature of +4°C under conditions of flowing Baikal water and a 12-hour light regime.

Analysis of biochemical parameters was carried out using the known methods: the pigment composition was analysed using a MALDI-TOF/TOF Mass Spectrometer (UltrafleXtreme, Bruker Daltonics GmbH, Germany) and MTP AnchorChip 400/384 T F MALDI target (S/N 20823) as well as by high-performance liquid chromatography (HPLC) on a Milikhrom A-02 chromatograph (EkoNova, Novosibirsk); gas chromatography-mass spectrometric analysis of methyl esters in fatty acids was carried out on an Agilent 5973N-GC6890 instrument.

Experiments on keeping *Lubomirskia baicalensis* in temperature-controlled aquarium installations have indicated that after three months of incubation of the sponge with the temperature rise from +4°C to +12°C quantitative biochemical parameters of cells in both the sponge itself and its symbionts change. The results of the experiment revealed that when the water temperature rises above the ecological optimum for the endemic species of the Baikal sponge, *Lubomirskia baicalensis*, the quantitative composition of demospongic acids will change. The cells of the sponge itself synthesize directly these acids, and they are taxonomic and trophic markers.

**Keywords**: Lake Baikal hydrobionts, Baikal sponges, lipids, fatty acids
the quantitative composition of demospongic acids will change. The cells of the sponge itself synthesize directly these acids, and they are taxonomic and trophic markers.

The present work was supported by the State Task No. 0345–2016–0002 (AAAA-A16-116122110066-1) “Molecular ecology and evolution of living systems in Central Asia under global environmental changes”; by RFBR and DST according to the research project № 19-54-45034 and RFBR № 20-04-00868; using unique facilities of “Experimental freshwater aquarium complex of Baikal hydrobionts” (LIN SB RAS).

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