#### **Short communication**

# Composition of physiological bacterial groups in thermal groundwaters of the Kuldur deposit (Amur oblast, Russian Far East)



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**ABSTRACT.** The research was focused on the distribution, composition, structure, numbers and diversity of predominant bacteria of different ecologo-trophic groups in the thermal groundwaters of the Kuldur deposit (Amur Oblast, the Russian Far East). The research results showed uneven distribution of bacteria of different physiological groups in the groundwaters as well as their low numbers. Thionic and iron-oxidizing bacteria predominated in all studied drill holes. Ammonium and nitrite-oxidizing bacteria were also numerous while *Bacillus* sp., *Paenibacillus* sp., *Brevibacillus* sp. were prevalent in the separated pure cultures

Keywords: Kuldur, thermal waters, bacteria, physiological groups, numbers

#### 1. Introduction

The Kuldur deposit of thermal waters is situated in the intracontinental southern part of the Russian Far East, near the settlement of Kuldur of the Jewish Autonomous Region. The Kuldur Springs crop out in the central part of Pionersky granite massif and are confined to the meridional fault at the intersection with feathering diastrophism of the north-east direction. Paleozoic granites and granodiorites are the enclosing rocks. Water temperature goes up to 73°C (Kompanichenko and Poturai, 2015). Favorable balneological properties allow using these thermal waters for therapeutic purposes in the resorts "Kuldur", "Voenny", "Sanus" and "Gornyak" that are situated in that area.

According to V.I. Vernadsky, water takes a special place among the chemical compounds of the Earth and natural groundwaters are complex dynamic systems that are closely interconnected with the surrounding environment (Vernadsky, 1965). The content and activity of microorganisms in groundwater has a significant influence on its geochemical parameters. The thermophiles that populate high-temperature environments play an important role in biogeochemical processes in ground biotopes. They transform subsurface rocks and gasohydrothermal chemical compounds by synthesizing organic matter and changing the composition of minerals that contain sulfur, iron, silicone and other chemical elements.

Activity of various functional groups of bacteria results in a rapid decrease in the migration intensity of some elements and leads to formation of minerals (Franke and Bazylinski, 2003; Romanenko and Karnachuk, 2004). Study of microorganisms in thermal habitats provides a better understanding of the origin and evolution of the early life and gives an opportunity to evaluate the patterns of their biogeochemical processes.

As the bacterial communities of the mineral waters in this region have not been numerous or sufficiently comprehensive (Fisher and Kompanichenko, 2007; Kalitina et al., 2017a; Lebedeva and Kharitonova, 2019) our research was aimed at studying the distribution, composition and numbers of different physiological bacterial groups in the thermal groundwaters of the Kuldur deposit.

### 2. Materials and methods

The objects of our research included thermal groundwaters of the Kuldur deposit drill holes No.1-87, No. 2-87, No. 3-87, No. 3-51, and No. 5-51. The highest-temperature waters (71–73°C) are accessed by drill holes 1-87 and 2-87 at the Kuldur River bend. Drill holes 1-87 and 2-87 have both the highest temperature and the highest output that allows considering this part of the deposit its central zone that coincides with the area of maximum unloading of thermal springs. Drill holes 3-87 and 3-51 on the surface are situated only 20-30 meters away from the central drill holes and are also

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included into the water crushing zone; however, the water temperature (55–60°C) and the output are much lower there. On the flanks there are drill holes 5-51 with water temperature range of 23-28°C. The water from drill holes No. 3-87, 3-51 and 5-51 is not used by the resorts because of its lower temperature. Only monitoring observations are performed at these closed drill holes with no water extraction; therefore their hydrodynamic pattern is quite stagnant. Nowadays, all drill holes at the Kuldur thermal field are suspended except for the five of them (1-87, 2-87, 3-87, 3-51, 5-51) (Poturai, 2013).

Thermal water samples were collected into sterile 1000 ml. tubes in triplicate in August, 2015. Prior to the analysis, all samples were stored in the refrigerator at 4° C for less than 24 hours. Traditional methods of practical microbiology were applied for revealing and cultivating the bacteria (Lysak et al., 2015). The quantity of different ecologo-trophic groups of bacteria was determined by the method of limiting dilution and the Koch's method (Egorov, 1995). Microorganisms were separated from groundwater with the method of membrane filtration through cellulose nitrate membrane filters with pore size of  $0.22~\mu m$ and 47 mm in diameter. Saprophytic bacteria were cultivated on organic-enriched agar (OA) and OA 1:10. Other physiological bacterial groups were identified in special selective media (Kuznetsov and Dubinina, 1989). Anaerobic bacteria of different physiological groups were cultivated in an aerostat using GasPak EZ gas generating sachets. Bacteria were cultivated in a thermostat at temperature 25°C (drill holes 5-51) and 60°C (drill holes 1-87, 2-87, 3-87, 3-51) within 14 days. Bacteria were identified with molecular-genetic methods.

# 3. Results and discussion

The Kuldur thermal spring is situated within the Khingan hydrogeological massif in the Amur-Okhotsk hydrogeological fold area. Water is recharged by infiltration of atmospheric precipitation and water inflows from the adjacent elevations; the drainage of water occurs through the local hydrographic network and the springs on valley slopes. Earlier surveys show that the Kuldur thermal waters are characterized by high temperatures (up to 73°C), low mineralization (< 1g/l), pH over 9 and the following chemical composition: HCO<sub>3</sub> (CO<sub>3</sub>)-Na and SO<sub>4</sub>-Na. The waters are substantially enriched in fluorine (up to 27 mg/l), silicone (up to 100 mg/l H<sub>2</sub>SiO<sub>2</sub> and over), aluminum (up to 10 mg/l and over), wolfram, molybdenum and some other anionogenic elements. The content of carbon total organic is insignificant and varies between 0.3–3.7 mg/dm<sup>3</sup>. Thermal water is colorless, bluish, and completely transparent with a slight smell of hydrogen sulfide. The total mineralization of water is 0.30-0.36 g/l, alkaline reaction (pH – 9.4- 9.7), radioactivity 1.2 – 2.4 Mache units. Both dissolved and spontaneous gases of thermal groundwaters of these drill holes mostly include nitrogen (up to 98 %) with admixed inert gases (Ar, Kr, Xe, He and Ne). Admixes pf other gases ( $CO_2$ ,  $H_2S$ ,  $O_2$ ) are, as a rule, insignificant (Poturai, 2013; Lyamina et al., 2019).

The results of microbiological analysis have shown that, in general, the groundwaters of the drill holes under study contain low numbers of cultivated physiological bacterial groups that, on average, comprised  $0.2 \times 10^2$  (1-87)  $-5.6 \times 10^3$  c/ml (5-51). Similarly low numbers of bacteria were earlier found by us in different geochemical types of groundwaters in the Russian Far East (Kalitina et al., 2017b; Kalitina et al., 2018; Lebedeva et al., 2019). The lowest composition of ecologo-trophic bacterial groups and their low quantity were found in higher-temperature groundwaters of central drill holes No.1-87 and No. 2-87 (Table) that corresponded to the minimum values of carbon total organic in those waters (Kompanichenko and Poturai, 2015). Aerobic and anaerobic saprophyte bacteria were not discovered in 100 ml water samples from groundwaters of drill holes No.1-87 and No.2-87. The lowest quantity of physiological groups was found in the water of drill hole No.1-87: only thionic bacteria were discovered there  $(2.8 \times 10^2 \text{ c/ml})$  thus showing the prevailing bacteria-induced oxidation of reduced sulfur compounds. In addition to thionic bacteria, a small amount of nitrite-oxidizing and anaerobic iron-reducing microorganisms were also discovered in drill holes No.2-87 (Table). Groundwaters of drill holes No.3-87, No.3-51 and No.5-51 were characterized with development of greater quantity of different physiological bacterial groups (10-12 out of 16) (Table); it can be attributed to the lower temperature of thermal waters (28-61°C) that is more favorable for bacterial activity. These waters also have a more stagnant hydrochemical water pattern that probably contributed to formation of specific microbiocenoses. The thermal waters of drill holes No.3-87, No.3-51, and No.5-51 showed prevalence of thionic and anaerobic iron-reducing bacteria, that corresponds to the previously obtained data (Kalitina et al., 2017a) and shows that microorganism-induced oxidation of reduced sulfur compounds and reduction of iron oxides are prevalent in the groundwaters. The quantity of predominant bacterial groups was lower in low-temperature waters (drill holes 5-51) whereas the quantity of ammonia-oxidizing and saprophyte anaerobic microorganism was the highest (Table) thus showing the processes of organic matter decomposition and oxidation of ammonia compounds. Predominant bacteria were separated from the thermal waters and identified. It was established that bacteria Bacillus sp., Paenibacillus sp., Brevibacillus sp. were significantly prevalent in the groundwaters of the drill holes under study.

## 4. Conclusions

Thus, the research shows that thermal groundwaters of the Kuldur drill holes are characterized with uneven distribution and low numbers of physiological bacterial groups. The hottest waters contained the lowest numbers and quantity

Dominant physiological groups of bacteria (cells/ml): 1-87 (72°C)		Groundwater Wells:				
		1-87 (72°C)	2-87 (72°C)	3-87 (61°C)	3-51 (60°C)	5-51 (28°C)
Saprophytes	aerobes	0	0	0	0	0
	anaerobes	0	0	$0.3 \times 10^{1}$	$0.5 \times 10^{1}$	$6.5 \times 10^{1}$
Thionic		$2.8 \times 10^{2}$	$4.3 \times 10^{2}$	$3.3 \times 10^{3}$	$2.4 \times 10^{2}$	$5.6 \times 10^{3}$
Sulfate Reducing		0	0	0	$0.7 \times 10^{2}$	0
Ammonium oxidizing		0	0	$1.4 \times 10^{1}$	$0.5 \times 10^{2}$	$3.2 \times 10^{2}$
Nitrite oxidizing		0	$0.5 \times 10^2$	$0.7 \times 10^{1}$	0	$0.5 \times 10^{2}$
Iron oxidizing autotrophs		0	0	0	$1.0 \times 10^{2}$	0
Iron reducing		0	$1.2 \times 10^{2}$	$5.1 \times 10^{2}$	$1.9 \times 10^{2}$	$2.8 \times 10^{2}$
The number of developing physiological groups of 16:		1	3	11	12	10
Average number		$0.2 \times 10^{2}$	$0.3 \times 10^{2}$	$2.4 \times 10^{2}$	$0.4 \times 10^{2}$	$3.9 \times 10^{2}$

Table. Distribution of dominant physiological groups of bacteria in the underground waters of the Kuldur deposit

of physiological bacterial groups. Thionic and ironreducing bacteria were predominant in all drill holes involved in this survey.

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