Effect of colchicine on physiological and biochemical properties of *Rhodococcus qingshengii*

Yu.A. Markova1, L.A. Belovezhets2, V.N. Nurminsky3, I.S. Kapustina4, N.V. Ozolina1, V.V. Gurina1, A.L. Rakevich5, A.V. Sidorov1

1 Siberian Institute of Plant Physiology and Biochemistry of Siberian Branch of Russian Academy of Sciences, Irkutsk, Russia
2 A.E. Favorsky Irkutsk Institute of Chemistry of the Siberian Branch of the Russian Academy of Sciences, Irkutsk, Russia
3 Irkutsk Branch of the Institute of Laser Physics, The Siberian Branch of the Russian Academy of Sciences, Irkutsk, Russia
4 juliam06@mail.ru

**Abstract.** The genus *Rhodococcus* includes polymorphic non-spore-forming gram-positive bacteria belonging to the class Actinobacteria. Together with *Mycobacterium* and *Corynebacterium*, *Rhodococcus* belongs to the Mycolata group. Due to their relatively high growth rate and ability to form biofilms, *Rhodococcus* are a convenient model for studying the effect of biologically active compounds on pathogenic *Mycobacteria*. Colchicine was previously found to reduce biofilm formation by *P. carotovorum* VKM B-1247 and *R. qingshengii* VKM Ac-2784D. To understand the mechanism of action of this alkaloid on the bacterial cell, we have studied the change in the fatty acid composition and microviscosity of the *R. qingshengii* VKM Ac-2784D membrane. Nystatin, which is known to reduce membrane microviscosity, is used as a positive control. It has been found that colchicine at concentrations of 0.01 and 0.03 g/l and nystatin (0.03 g/l) have no significant effect on the survival of *R. qingshengii* VKM Ac-2784D cultivated in a buffered saline solution with 0.5 % glucose (GBSS). However, colchicine (0.03 g/l) significantly inhibits biofilm formation. *Rhodococcus* cells cultivated for 24 hours in GBSS with colchicine acquire a rounded shape. Colchicine at 0.01 g/l concentration increases C16:1(n-7), C17:0, C20:1(n-9) and C21:0 fatty acids. The microviscosity of the membrane of individual cells was distributed from the lowest to the highest values of the generalized laurdan fluorescence polarization index (GP), which indicates a variety of adaptive responses to this alkaloid. At a higher concentration of colchicine (0.03 g/l) in the membranes of *R. qingshengii* VKM Ac-2784D cells, the content of saturated fatty acids increases and the content of branched fatty acids decreases. This contributes to an increase in membrane microviscosity, which is confirmed by the data on the GP fluorescence of laurdan. All of the above indicates that colchicine induces a rearrangement of the *Rhodococcus* cell membrane, probably in the direction of increasing its microviscosity. This may be one of the reasons for the negative effect of colchicine on the formation of *R. qingshengii* VKM Ac-2784D biofilms.

Key words: *Rhodococcus qingshengii*; colchicine; biofilms; fatty acids; membrane microviscosity.

For citation: Markova Yu.A., Belovezhets L.A., Nurminsky V.N., Kapustina I.S., Ozolina N.V., Gurina V.V., Rakevich A.L., Sidorov A.V. Effect of colchicine on physiological and biochemical properties of *Rhodococcus qingshengii*. *Vavilovskii Zhurnal Genetiki i Selektsi = Vavilov Journal of Genetics and Breeding*. 2022;26(6):568-574. DOI 10.18699/VJGB-22-69

Влияние колхицина на физиолого-bióхимические свойства *Rhodococcus qingshengii*

Ю.А. Маркова1, А.А. Беловежеч2, В.Н. Нурминский1, И.С. Капустина1, Н.В. Озолина1, В.В. Гурина1, А.А. Ракевич3, А.В. Сидоров1

1 Сибирский институт физиологии и биохимии растений Сибирского отделения Российской академии наук, Иркутск, Россия
2 Иркутский институт химии им. А.Е. Фаворского Сибирского отделения Российской академии наук, Иркутск, Россия
3 Иркутский филиал Института лазерной физики Сибирского отделения Российской академии наук, Иркутск, Россия
4 juliam06@mail.ru

**Аннотация.** Род *Rhodococcus* объединяет полиморфные неспоробразующие грамположительные бактерии, относящиеся к классу Actinobacteria. *Rhodococcus* вместе с *Mycobacterium* и *Corynebacterium* входят в группу *Mycobacteria*. Благодаря относительно высокой скорости роста и способности к образованию биопленок, *Rhodococcus* являются удобной моделью для изучения действия биологически активных соединений на патогенные *Mycobacteria*. Ранее было показано, что колхицин угнетал образование биопленок у *P. carotovorum* VKM B-1247 и *R. qingshengii* VKM Ac-2784D. Целью настоящей работы было изучение действия колхицина на жизнеспособность и микровязкость мембран *Rhodococcus qingshengii* VKM Ac-2784D для понимания механизма действия этого алкалоида на бактериальную клетку. В качестве положительного контроля использовали нистатин, снижающий микровязкость мембран. Установлено, что колхицин в концентрациях 0.01 и 0.03 г/л и нистатин (0.03 г/л) не оказали существенного влияния на выживаемость *R. qingshengii* VKM Ac-2784D, культивируемого в buffered saline solution с 0.5 % glucose (GBSS). Однако колхицин (0.03 г/л) существенно ингибирует образование биопленок. *Rhodococcus* в течение 24 часов в GBSS с колхицином приобретает округлую форму. Колхицин при концентрации 0.01 г/л увеличивает содержание C16:1(n-7), C17:0, C20:1(n-9) и C21:0 жировых кислот. Микровязкость мембраны клеток индивидуальных *Rhodococcus* была распределена от наименьших до наибольших значений индекса поляризации поляризованного излучения laurdan (GP), что указывает на большой набор адаптивных реакций к данному алкалоиду. При более высокой концентрации колхицина (0.03 г/л) в мембранах *R. qingshengii* VKM Ac-2784D клеток, содержание насыщенных жировых кислот увеличивается и содержание жировых кислот с ветвями уменьшается. Это приводит к увеличению микровязкости мембраны, что подтверждается данными по индексу поляризации laurdan. Все описанные выше свидетельствуют, что колхицин индуцирует перестройку мембраны *Rhodococcus*, вероятно в направлении увеличения ее микровязкости. Это может быть одной из причин негативного действия колхицина при формировании биопленок *R. qingshengii* VKM Ac-2784D.

Ключевые слова: *Rhodococcus qingshengii*; колхицин; биопленки; жировые кислоты; микровязкость мембраны.

Для цитирования: Маркова Ю.А., Беловежеч А.А., Нурминский В.Н., Капустина И.С., Озолина Н.В., Гурина В.В., Ракевич А.А., Сидоров А.В. Влияние колхицина на физиолого-биохимические свойства *Rhodococcus qingshengii*. *Vavilovskii Zhurnal Genetiki i Selektsi = Vavilov Journal of Genetics and Breeding*. 2022;26(6):568-574. DOI 10.18699/VJGB-22-69
**Introduction**

The *Rhodococcus* genus includes polymorphic non-spore-forming gram-positive bacteria that belong to the Actinobacteria class. *Rhodococcus* are frequently met in nature, in particular, in living organisms. Among the key features of these microorganisms is their ability to decompose different organic compounds, including pollutants (PAHs, biphenyls, alkanes, etc.) (Szőköl et al., 2014; Li et al., 2016). For this reason, *Rhodococcus* continue to attract the growing interest as valuable biotech species.

Along with *Mycobacterium* and *Corynebacterium*, *Rhodococcus* relates to the Mycolata group, which is characterized by the presence of mycolic acids on the cell walls (Sutcliffe, 1998). This makes these bacteria more resistant to the toxic compounds such as disinfectants, antibiotics or PAC. Unlike myco- and corynebacteria, *Rhodococcus* species are mostly non-pathogenic. Therefore, owing to relatively high growth rate and propensity to biofilm formation, the *Rhodococcus* represent a convenient model to examine the effect of biologically active compounds on pathogenic *Mycobacteria*.

The integrity of a microbial cell drastically depends on the membrane. In order to survive in ever-changing environmental conditions and to maintain optimal membrane fluidity, the bacteria change the fatty acid composition of membrane lipids (Dubois-Brissonnet et al., 2016). The cell membrane is the major target of non-polar organic solvent toxicity (De Carvalho et al., 2005). Plant metabolites also affect the membrane via inhibition of the efflux channels activity (Tegos et al., 2002), the content of porin proteins (Abreu et al., 2012), etc.

Previously, we found that the alkaloid colchicine at a concentration of 0.25 g/l suppressed the formation of a biofilm by *Pectobacterium carotovorum* VKM B-1247 and *Rhodococcus qingshengii* VKM Ac-2784D species (Bybin et al., 2018). Moreover, no negative effect on the viability of these bacteria was revealed. Colchicine is widely known as an alkaloid that interrupts the tubulin polymerization in eukaryotic cells (Zhang et al., 2018). It is likely that colchicine exhibits a similar effect on the microorganisms, affecting the cytoskeleton and preventing the adhesion of microbial cells (Dubey et al., 2011). However, its influence on microbial cells was poorly studied. All of the above sparked our interest in this compound.

In the present work, we have examined the effect of colchicine on the fatty acid composition and microviscosity of *R. qingshengii* VKM Ac-2784D membranes.

**Materials and methods**

*R. qingshengii* VKM Ac-2784D strain isolated from the rhizosphere of couch grass (*Elytrigia repens* (L.) Nevski) growing in the oil-contaminated territory of the Irkutsk region (Russia) was used in the work (Petrushin et al., 2021). The *Rhodococcus* strain features a good formation of biofilms, and therefore represents a convenient model for their study.

The bacteria were cultivated on BTA-agar (Biotekhnovatsiya, Russia) for 48 h at 26 °C. Then they were transferred to a 0.5 % glucose buffered saline solution (GBSS) and the density of the suspension was adjusted to OD$_{595}$ 0.26–0.33.

The minimum inhibitory concentration (MIC) of colchicine for *R. qingshengii* VKM Ac-2784D was determined by the limiting dilution method (Guidelines..., 2000).

To evaluate the effect of nystatin and colchicine on growth kinetics and biofilm formation, 150 μl of bacterial suspension was added to the wells of sterile flat-bottom 96-well plates and the optical density was measured on the first, third, and eighth days of cultivation using an iMark plate reader (Bio-Rad, USA), $\lambda = 595$ nm. The plate was washed from loosely attached cells. The precipitate was stained with 1 % crystal violet solution for 45 min. After washing (3 times) to extract the dye, 200 μl of 96 % ethanol was added to the wells. The level of extraction (absorption) of crystalline violet with ethanol was measured using an iMark plate reader (Bio-Rad) at a wavelength of 595 nm in optical density units (OD$_{595}$). The degree of biofilm formation corresponded to the intensity of dye staining of the wells content (Shaginyan et al., 2007).

Two controls were employed in the work. The first one involved the bacteria cultivated in GBSS without the addition of colchicine. The second control used the bacteria grown in a medium with 0.03 g/l of nystatin (Biosintez, Russia), since nystatin can reduce microviscosity of the cell membranes. Colchicine (Sigma-Aldrich, USA) was applied at concentrations of 0.01 and 0.03 g/l. When plotting the diagrams, the relative optical density in % to the control was used. Cell sizes were assessed using the AxioVision Rel 4.8 software.
To determine the fatty acid composition of the bacterial membrane and the orderliness (microviscosity, fluidity) of its lipid phase, the bacteria were cultivated in the aforementioned media for a day. The membrane lipids orderliness was evaluated by the generalized polarization (GP) of laurdan lipophilic probe fluorescence in each pixel corresponding to the luminescent image domain. To stain the bacteria, 10 μM of a methanolic solution of laurdan (2-(dimethylamino)-6-dodecanoylnapthalene) (Sigma-Aldrich) was added to each vial. Live stained bacteria were observed using a microscope (laser scanning confocal fluorescent microscope MicroTime 200; PicoQuant GmbH, Germany).

The distribution of GP values was analyzed by visualization with histograms. For each histogram, a theoretical multimodal distribution as a superposition of several normal distributions was plotted (Nurminsky et al., 2017). Next, the parameter fitting of the experimental distributions of bacterial membrane GP values was estimated. The model distribution was a normal distribution or a mixture of distributions and thus consisted of one or more components. Finally, the optimal parameters of the components that were closest to the experimental distribution were selected.

To determine the composition of fatty acids (FA), the bacteria were cultivated similarly without the addition of laurdan. The lipids were extracted according to the published procedure (Bligh, Dyer, 1959). After removal of the solvent, a 1 % methanol solution of H₂SO₄ was added to the lipid extract and heated on a water bath at 60 °C for 30 min. After cooling, the solution was extracted (3 times) with hexane (Christie, 1993). Fatty acids methyl esters were analyzed using an Agilent technology 5973N/6890N MSD/DS chromato-mass spectrometer (USA). Detector (mass spectrometer) was quadrupole, ionization method was electron impact (EI), ionization energy was 70 eV, the mode of the total ion current registration was used for the analysis. Separation was performed on an HP-INNOWAX capillary column (30 m × 250 μm × 0.50 μm). The stationary phase was polyethylene glycol. The mobile phase was helium; gas flow rate was 1 ml/min. Temperature of the evaporator was 250 °C, temperature of the ion source was 230 °C, temperature of the detector was 150 °C, and temperature of the line connecting the chromatograph with the mass spectrometer was 280 °C. Scan range was 41–450 amu. The volume of the injected sample was 1 μl, the flow separation was 5:1. Chromatography was carried out in isothermal mode at 200 °C. To identify the peaks of FA methyl esters, methyl ester standards (Sigma-Aldrich) and mass spectrometry using the NIST 05 mass spectrum library (Ozolina et al., 2017) were used. The content of individual fatty acids was calculated as a percentage of the total amount of fatty acids and divided into groups: saturated (SFA), monounsaturated (MUFA), polyunsaturated (PUFA), saturated iso- and anteiso-methyl branched fatty acids (BFA) (Rodrigues, de Carvalho, 2015).

The significance of differences in biofilm formation and the quantitative content of fatty acids were assessed using the nonparametric Kruskal–Wallis test with Dunnett’s correction (Glantz, 1991). All calculations were performed using the RStudio software.

Results and discussion
It was found that the MIC of colchicine for *R. qingshengii* VKM Ac-2784D is 0.02 g/l. Therefore, in further experiments, the concentrations below and above this value, i.e. 0.01 and 0.03 g/l, respectively, were used. The selected concentrations of colchicine and nystatin did not strongly affect the growth of *Rhodococcus* (Fig. 1). At the same time, it was established that colchicine at a concentration of 0.03 g/l significantly inhibited the formation of a biofilm at all stages of the experiment, while at a concentration of 0.01 g/l a divergent effect was observed. On the first day, nystatin stimulated the formation of a biofilm, whilst on the third and eighth days of cultivation, its effect was comparable to the control (Fig. 2).

![Fig. 1. Optical density of *R. qingshengii* VKM Ac-2784D cell suspension relative to control, %, on the first, third and eighth days of cultivation.](image1)

Here and in the Fig. 2: 1 – GBSS; 2 – GBSS with 0.03 g/l colchicine; 3 – GBSS with 0.01 g/l colchicine; 4 – GBSS with 0.03 g/l nystatin.

![Fig. 2. Optical density of biofilm *R. qingshengii* VKM Ac-2784D on the first, third and eighth days of cultivation.](image2)
The cultivation of *R. qingshengii* VKM Ac-2784D in the presence of colchicine for a day essentially changed the cell morphology: the cells acquired a more rounded shape (Table 1). Moreover, the intracellular content became heterogeneous (Fig. 3), which is consistent with the results obtained for *Bacillus megaterium* (Dubey et al., 2011). The shape of *Rhodococcus* cells under the action of nystatin remained intact.

The changes in cell morphology are usually accompanied by structural and functional rearrangement of their cell membranes (de Carvalho et al., 2014). Specifically, the degree of saturation of FA, their length, as well as the amount of branched fatty acids are altered.

Under the control conditions, the *Rhodococcus* cell membranes mainly contained palmitic, stearic, and oleic acids (Table 2). The ratio of saturated to monounsaturated fatty acids was 1.64. The content of polyunsaturated and branched acids was small (4.41 and 0.84 %, respectively). Colchicine at a concentration of 0.01 g/l changed the ratio of saturated and monounsaturated FA (1.29) in favor of the latter, and simultaneously reduced the amount of polyunsaturated FA. At the same time, the number of long-chain FA C20:1(n-9), C21:0 and C22:0 increased. With 0.03 g/l of colchicine, the relative content of saturated and branched FA increased (Fig. 4), while the ratio of UFA to MUFA reached 1.89. All this indicates the rearrangement of *R. qingshengii* VKM Ac-2784D membrane after the introduction of colchicine into the cultivation medium. Interestingly, different concentrations of colchicine had an opposite effect on the composition of cell membrane FA. This is probably due to various degrees of regulatory systems disorder. The addition of nystatin, a compound that fairly increases the membrane fluidity, led to a higher content of unsaturated and branched FA, substantially lower amount of palmitic FA, higher concentration of oleic FA.

The fluidity or microviscosity of membranes is an integral index that depends on lipid saturation and content of sterols or proteins. Therefore, further we focused our efforts on the evaluation of the colchicine and nystatin effect on the orderliness of the lipid phase of *R. qingshengii* VKM Ac-2784D membrane. For this purpose, the laurdan fluorescence GP index was used, which can vary from –1 to +1. Its negative values correspond to lower microviscosity (higherfluidity) of the cell membrane (Nurminsky et al., 2015) (see Materials and Methods).

The fitting of experimental distributions of bacterial membrane GP values permitted to find from one to four components under the action of nystatin and colchicine (Fig. 5). In all variants, the most significant component characterizes the liquid-disordered regions of the membrane (α (average GP values): –0.16–0.04, contribution: 73.9–100 %). The sterol-binding agent nystatin shifted GP towards a decrease in the orderliness of the membranes (α: –0.16, contribution: 100 %), which corresponds to the known mechanism of action of this antibiotic on the membranes of eu- and prokaryotes (Efimova et al., 2014). Colchicine, on the contrary, increased the orderliness of the membranes: a of the most significant components shifted, although slightly, towards positive values compared to the control in both concentrations (α: 0.04, contribution: 73.9–89.4 %), which agrees with the observed increase in the amount of saturated FA. However, this significantly expanded the data scattering, and the number of components reached 2 (in the variant with

---

**Table 1.** Cell sizes of *R. qingshengii* VKM Ac-2784D after cultivation for a day under control conditions and in the presence of the studied compounds

| Experiment          | Length, nm | Width, nm | Length/width |
|---------------------|------------|-----------|--------------|
| Control             | 1.35 ± 0.29| 0.56 ± 0.08| 2.42         |
| Colchicine, 0.01 g/l| 1.04 ± 0.23*| 0.58 ± 0.1| 1.78*        |
| Nystatin, 0.03 g/l  | 1.46 ± 0.36| 0.68 ± 0.14| 2.12         |

* p < 0.05.

---

**Fig. 3.** Cell morphology of *R. qingshengii* VKM Ac-2784D, cultivated for 24 hours in buffered saline with glucose (5 g/L) (C) and after adding 0.03 g/l nystatin (1), 0.01 g/l colchicine (2), 0.03 g/l colchicine (3).

Stained with laurdan. Magnification × 600.
Effect of colchicine on physiological and biochemical properties of *Rhodococcus qingshengii*

**Table 2.** The composition of FA in the membrane cells of *R. qingshengii* VKM Ac-2784D cultivated for a day in GBSS with colchicine (0.03 and 0.01 g/l) and nystatin (0.03 g/l)

| Fatty acid          | Control               | Concentration of alkaloids, g/l |
|---------------------|-----------------------|---------------------------------|
|                     |                       | Colchicine | Nystatin |
|                     |                       | 0.03       | 0.01     | 0.03     |
| Lauric C12:0        | –                     | –          | –        | 0.76 ± 0.04 |
| Myristic C14:0      | 4.76 ± 0.38           | 6.09 ± 0.02 | 5.45 ± 0.26 | 4.89 ± 0.02 |
| Pentadecane C15:0   | 1.99 ± 0.22           | 2.36 ± 0.32 | 2.25 ± 0.01 | 1.60 ± 0.06 |
| Anteiso-pentadecane C15:0-a | –           | –          | –        | 1.08 ± 0.03 |
| Palmitic C16:0      | 36.76 ± 3.52          | 34.58 ± 0.33 | 32.27 ± 2.10 | 27.98 ± 0.48* |
| Isopalmmitic C16:0-i| –                     | –          | –        | 1.63 ± 0.04 |
| Palmitoleic C16:1(n-9) | 5.16 ± 1.05           | 3.78 ± 0.35 | 4.50 ± 0.14 | 3.53 ± 0.21 |
| C16:1(n-7)          | 1.31 ± 0.14           | 2.33 ± 0.35 | 3.11 ± 0.36* | 2.21 ± 0.28 |
| C16:1(n-9)          | –                     | –          | –        | 4.78 ± 0.17 |
| C16:1(n-5)          | 2.30 ± 0.26           | 2.06 ± 0.09 | 3.79 ± 1.04 | –        |
| Heptadecane C17:0   | 1.26 ± 0.14           | 1.35 ± 0.03 | 1.60 ± 0.11* | 1.28 ± 0.05 |
| Anteiso-heptadecane C17:0-a | 0.84 ± 0.23           | 1.23 ± 0.03* | 0.72 ± 0.16 | 0.83 ± 0.09 |
| Stearic C18:0       | 12.09 ± 0.19          | 14.80 ± 0.25 | 10.15 ± 0.21 | 13.91 ± 0.29 |
| Oleic C18:1(n-9)    | 25.79 ± 2.42          | 24.24 ± 0.24 | 28.46 ± 1.10 | 30.29 ± 0.09* |
| C18:1(n-7)          | 1.36 ± 0.11           | 0.89 ± 0.02 | 2.33 ± 0.13 | 0.59 ± 0.13 |
| Linolic C18:2(n-6)  | 4.41 ± 0.61           | 1.54 ± 0.01 | 2.01 ± 0.49 | 2.33 ± 0.05 |
| Arachic C20:0       | 0.99 ± 0.08           | 1.55 ± 0.01* | 1.16 ± 0.01 | 0.86 ± 0.00 |
| C20:1(n-9)          | –                     | 0.46 ± 0.04 | 0.36 ± 0.01 | 0.27 ± 0.00 |
| Heneicosanic C21:0  | –                     | 0.48 ± 0.09 | 0.44 ± 0.09 | 0.26 ± 0.03 |
| Behenic C22:0       | 1.10 ± 0.13           | 2.28 ± 0.19 | 1.40 ± 0.03 | 0.92 ± 0.17 |

* p < 0.05.

**Fig. 4.** Relative content of the main groups of fatty acids during cultivation for a day in buffered saline with glucose (5 g/l) (C) and with the introduction of 0.03 g/l nystatin (1), 0.01 g/l colchicine (2), 0.03 g/l colchicine (3).

SFA – saturated fatty acids; PUFA – polyunsaturated fatty acids; MUFA – mono-unsaturated fatty acids; BFA – saturated iso- and anteiso-methyl branched fatty acids.

0.01 mg/ml) and 4 (in the variant with 0.03 mg/ml). Minor components corresponded to more densely packed regions of the membranes (α: 0.29, contribution: 1.8 %) or, conversely, to less densely packed ones (α: –0.29, contribution: 6.7 %).

**Conclusion**

In conclusion, colchicine in the composition of GBSS at concentrations of 0.01 and 0.03 g/l did not significantly affect the survival of *R. qingshengii* VKM Ac-2784D, but strongly inhibited the formation of a biofilm. *Rhodococcus* cells cultivated for 24 hours in GBSS with colchicine acquired a rounded shape. With 0.01 g/l of colchicine, the content of C16:1(n-7), C17:0, C20:1(n-9) and C21:0 FA acids increased.

The membrane microviscosity of individual cells is distributed from the lowest to the highest GP values, which indicates a variety of adaptive responses to this alkaloid. At a higher concentration of colchicine (0.03 g/l) in the cell membranes of *R. qingshengii* VKM Ac-2784D, the
content of saturated fatty acids increased, while the amount of branched fatty acids reduced. This enhanced the membrane microviscosity that was confirmed by the values of laurdan fluorescence GP. These data testify to an adaptive rearrangement of the cell membrane under the action of the studied alkaloid, which is consistent with the results obtained by other authors (Wang et al., 2020). This may be a reason of the negative effect of colchicine on the formation of *R. qingshengii* VKM Ac-2784D biofilms.

**References**

Abreu A.C., McBain A.J., Simoes M. Plants as sources of new antimicrobials and resistance-modifying agents. *Nat. Prod. Rep.* 2012; 29(9):1007-1021. DOI 10.1039/c2np20035j.

Bligh E.G., Dyer W.J. A rapid method of total lipid extraction and purification. *Can. J. Biochem. Physiol.* 1959;37:911-917.

Bybin V.A., Turskaya A.L., Maksimova L.A., Markova Yu.A. Evaluation of the influence of some alkaloids on biofilm formation by different bacterial species. In: Proceedings of the Annual Meeting of the Society of Plant Physiologists of Russia, the All-Russia Scientific Conference with International Participation, and the School of Young Scientists. Irkutsk: Sochava Institute of Geography of the Siberian Branch of the Russian Academy of Sciences, 2018;1206-1209. DOI 10.31255/978-5-94797-319-8-1206-1209. (in Russian)

Christie W.W. Preparation of ester derivatives of fatty acids for chromatographic analysis. In: Christie W.W. Advances in Lipid Methodology – Two. Dundee: Oily Press, 1993;69-111.

de Carvalho C.C., Marques M.P., Hachicho N., Heipieper H.J. Rapid adaptation of *Rhodococcus erythropolis* cells to salt stress by synthesizing polyunsaturated fatty acids. *Appl. Microbiol. Biotechnol.* 2014;98(12):5599-5606. DOI 10.1007/s00253-014-5549-2.

de Carvalho C.C., Parreño-Marchante B., Neumann G., Da Fonseca M.M.R., Heipieper H.J. Adaptation of *Rhodococcus erythropolis* DCL14 to growth on n-alkanes, alcohols and terpenes. *Appl. Microbiol. Biotechnol.* 2005;67(3):383-388. DOI 10.1007/s00253-004-1750-z.

Dubey K.K., Jawed A., Haque S. Structural and metabolic correlation for *Bacillus megaterium* ACBT03 in response to biotransformation. *Microbiology.* 2011;90(6):758-767. DOI 10.1134/ S0026261711060899.

Dubois-Brissonnet F., Trotier E., Briandet R. The biofilm lifestyle involves an increase in bacterial membrane saturated fatty acids. *Front. Microbiol.* 2016;7:1673. DOI 10.3389/fmicb.2016.01673.

Efimova S.S., Schagina L.V., Ostrounova O.S. Investigation of channel-forming activity of polyene macrolide antibiotics in planar lipid bilayers in the presence of dipole modifiers. *Acta Naturae.* 2014; 6(4(23)):67-79. DOI 10.32607/20758251-2014-6-4-67-79.

Glantz S. Primer of Biostatistics. McGraw-Hill Publ., 1991.

Guidelines for the Experimental (Preclinical) Study of New Pharmacological Substances. Moscow: Remedium Publ., 2000. (in Russian)

Li C., Zhang C., Song G., Liu H., Sheng G., Ding Z., Wang Z., Sun Y., Xu Y., Chen J. Characterization of a protocatechuate catabolic gene cluster in *Rhodococcus ruber* OA1 involved in napthalene degradation. *Ann. Microbiol.* 2016;66(1):469-478. DOI 10.1007/s13213-015-1132-z.

Nurminsky V.N., Nesterkina I.S., Spiridonova E.V., Ozolina N.V., Rakevich A.L. Identification of sterol-containing domains in vacuolar membranes by confocal microscopy. *Biochemistry (Moscow).* Supplement Series A: Membrane and Cell Biology. 2017;11(4):296-300. DOI 10.1134/S1990748717040080.

Nurminsky V.N., Ozolina N.V., Nesterkina I.S., Kolesnikova E.V., Salyaev R.K., Rakevich A.L., Martynovich E.F., Pilipchenko A.A., Chernyshov M.Y. Peculiar properties of some components in a plant cell vacuole morphological structure revealed by confocal microscopy. *Cell Tiss. Biol.* 2015;9(5):406-414. DOI 10.1134/S1990519115050090.

Ozolina N.V., Gurina V.V., Nesterkina I.S., Dudareva L.V., Katysheva A.I., Nurminsky V.N. Fatty acid composition of total lipids in vacuolar membrane under abiotic stress. *Biologicheskiye Membrany = Biological Membranes.* 2017;34(1):63-69. DOI 10.7868/S023 3475517010078. (in Russian)

Petrushin I.S., Markova Yu.A., Karpnova M.S., Zaytseva Yu.V., Belovzhezts I.A. Complete genome sequence of *Rhodococcus qingshengii* strain VKM Ac-2784D, isolated from *Elytrigia repens* rhizosphere. *Microbiol. Resour. Announc.* 2021;10(11):e00107-21. DOI 10.1128/MRA.00107-21.

Rodrigues C.J.C., de Carvalho C.C.C.R. *Rhodococcus erythropolis* cells adapt their fatty acid composition during biofilm formation on metallic and non-metallic surfaces. *FEMS Microbiol. Ecol.* 2015;91(12):fiw135. DOI 10.1093/femsec/fiw135.
Effect of colchicine on physiological and biochemical properties of *Rhodococcus qingshengii*

complex in dependence of their phenotypic and genotypic characteristics. *Zhurnal Mikrobiologii, Epidemiologii i Immunobiologii = Journal of Microbiology, Epidemiology, and Immunobiology.* 2007; 1:3-9. (in Russian)

Sutcliffe I.C. Cell envelope composition and organisation in the genus *Rhodococcus*. *Antonie Van Leeuwenhoek.* 1998;74(1):49-58. DOI 10.1023/a:1001747726820.

Szóköl J., Rucká L., Šimčíková M., Halada P., Nešvera J., Pátek M. Induction and carbon catabolite repression of phenol degradation genes in *Rhodococcus erythropolis* and *Rhodococcus jostii*. *Appl. Microbiol. Biotechnol.* 2014;98(19):8267-8279. DOI 10.1007/s00253-014-5881-6.

Acknowledgements. The work was carried out within the framework of the basic theme (Registration No. 121031300011-7).

Conflict of interest. The authors declare no conflict of interest.

Received May 18, 2022. Revised June 29, 2022. Accepted July 8, 2022.