

Original Research

# Specificity of *Rhodococcus opacus* 1CP cells' responses to benzoate and 3-chlorobenzoate

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## Abstract

**Background:** Halogenated aromatic compounds are more resistant to microbial degradation than non-halogenated aromatic compounds. Microbial degradation of sodium benzoate in the presence of sodium 3-chlorobenzoate is of interest. The ability to degrade aromatic compounds is largely determined by the substrate specificity of the first enzyme that initiates degradation, namely, benzoate 1,2-dioxygenase for benzoate degradation, and 3-chlorobenzoate 1,2-dioxygenase for 3-chlorobenzoate degradation. In this study, the perspective of immobilized cells of *Rhodococcus opacus* 1CP actinobacterium for degradation of benzoate and 3-chlorobenzoate was explored. **Methods:** The biosensor approach (a membrane microbial sensor based on immobilized cells of *Rhodococcus opacus* 1CP and the Clark-type oxygen electrode as a transducer) was applied to evaluate the actinobacterial cells' responses to benzoate and 3-chlorobenzoate in the absence of both enzymes, benzoate 1,2-dioxygenase and 3-chlorobenzoate 1,2-dioxygenase, or in the presence of one of the said enzymes. **Results:** Data obtained show that 1CP actinobacterium possessed a constitutive system for the transport of benzoate and 3-chlorobenzoate into culture cells. The affinity of the transport system for benzoate was higher than that for 3-chlorobenzoate. Moreover, adaptation to one substrate did not preclude the use of the second substrate. Probably, porins facilitated the penetration of benzoate and 3-chlorobenzoate into 1CP cells. Analyzing  $V$  vs.  $S$  dependencies, negative cooperativity was found, when benzoate 1,2-dioxygenase bound substrate (3-chlorobenzoate), while positive cooperativity was determined at benzoate binding. The observed difference could be associated with the presence of at least two systems of 3-chlorobenzoate transport into actinobacterial cells and allosteric interaction of active sites of benzoate 1,2-dioxygenase in the presence of 3-chlorobenzoate. **Conclusions:** The membrane microbial sensor based on immobilized *Rhodococcus opacus* 1CP cells could be useful as a perspective tool for comparative evaluation of enzymes of complex structure such as benzoate- and 3-chlorobenzoate 1,2-dioxygenase.

**Keywords:** *Rhodococcus opacus* 1CP actinobacterium; benzoate 1,2-dioxygenase; 3-chlorobenzoate 1,2-dioxygenases; immobilized cells; membrane microbial sensor

## 1. Introduction

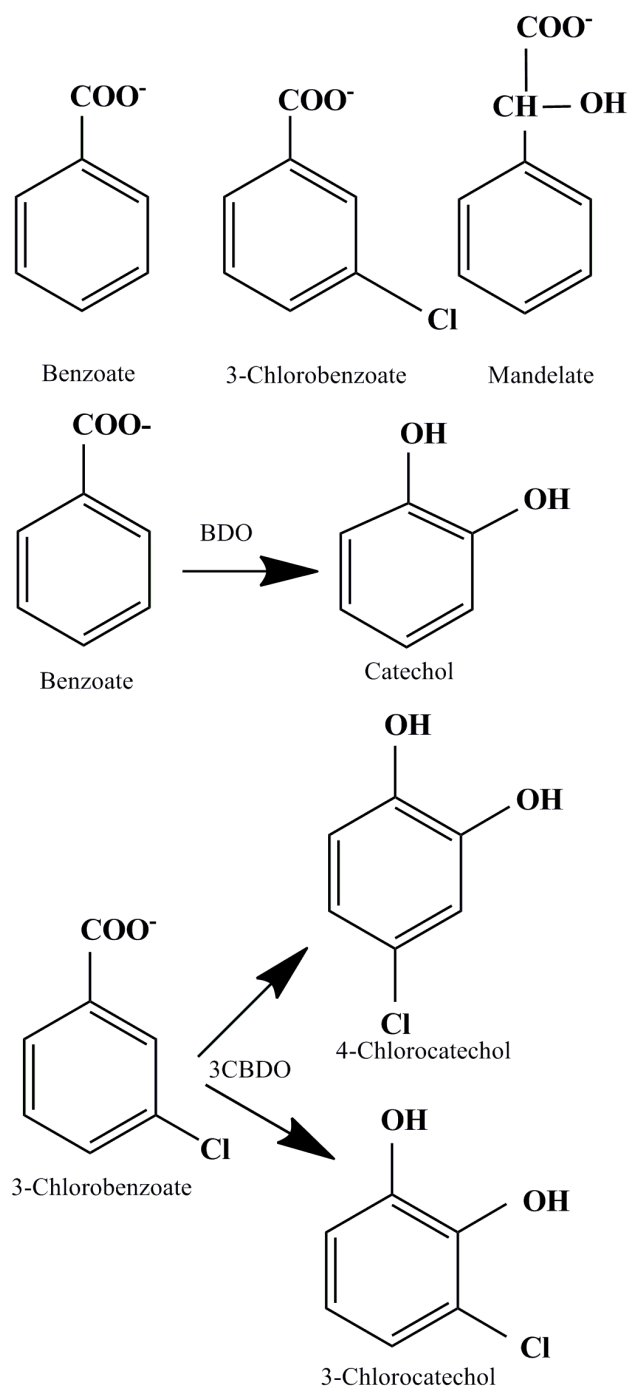
Salts of benzoic acid (benzoates) and chlorobenzoic acids (CBAs) enter the biosphere because of human activity in a household environment as well as during vital activity of living organisms and different reactions that occur in nature [1–9]. The harmfulness of benzoates is still discussed, but most of chlorinated aromatic compounds are toxic. It is known, that halogenated aromatic compounds are usually more resistant to microbial degradation than non-halogenated aromatic compounds. With an increase in the number of substituents in the ring, this resistance enhances. So, new remediation methods, including biological techniques, are being developed [2].

For bioremediation of polluted soil, two approaches such as activation of detoxifying ability of soil microflora and introduction of xenobiotics-degrading strains are used. Many species of the genus *Pseudomonas* and *Rhodococcus* can degrade a wide spectrum of xenobiotics [4,10–13]. The search for and introduction of new promising

strains able to degrade various pollutants is urgent. For example, the Gram-positive actinobacterium *Rhodococcus opacus* 1CP can be used to detoxify compounds such as (chloro)phenols, (chloro/methyl) benzoates including 3-chlorobenzoate (3CBA) [14,15]. To use a culture as a degrader of xenobiotics, it is important to know about the influence of xenobiotics on the microorganism and the activity of the enzymes involved in xenobiotic metabolism.

The ability to degrade (halogen-substituted) aromatic compounds is mainly determined by the substrate specificity of the first enzyme that initiates degradation. Benzoate 1,2-dioxygenase initiates benzoate metabolism (Fig. 1). It is considered that benzoate 1,2-dioxygenase is unable to perform an oxygenase attack on the 3-chlorobenzoate ring. Regarding the microbial degradation of 3CBA, there are several ways of degradation of this xenobiotic in bacterial cells. One of the ways is oxidation of the aromatic ring by 3-chlorobenzoate 1,2-dioxygenase enzyme [16] as shown in Fig. 1.





**Fig. 1.** Comparison of structural formulas of BA, 3CBA and mandelate; the first steps of BA and 3CBA metabolism in *R. opacus* ICP.

Benzoate 1,2-dioxygenase (BDO) is a two-component enzyme of complex structure that catalyzes the oxygen-dependent reaction [17]. 3-chlorobenzoate 1,2-dioxygenase (3CBDO) is the enzyme of peripheral metabolism that initiates the degradation of 3CBA. This enzyme catalyzes the reaction with the involvement of oxygen. Although activity of stable soluble enzymes may be determined in cell-free extracts of culture, activities

of BDO and 3CBDO cannot be determined in cell-free extracts because the enzymes are destroyed during preparation of cell-free extracts [18]. For that reason, BDO and 3CBDO activities are determined in whole cells by a change of oxygen consumption by cells in the presence of the substrate of the enzyme under study [19–21].

The biosensor on the basis of bacterial cells is a helpful convenient tool for rapid assessment of the substrate specificity and activity of the enzyme initiating the pathway in this bacterium [22]. A whole-cell biosensor with the Clark-type oxygen electrode as a transducer can be used for estimation of a change in oxygen consumption by microbial cells under the action of a substrate. In our previous study, BDO and 3CBDO activities in *R. opacus* ICP cells was determined by means of a biosensor method with the use of both a suspension of freshly harvested intact cells and immobilized resting cells [23,24]. The advantages of the whole-cell biosensor method for determination of enzyme activity are small amounts of microbial biomass for biosensor system formation and rapidness of assay.

Besides the measurement of the enzyme activity, whole cell biosensors can be used for detection of toxic aromatic compounds. To improve selectivity of a microbial sensor, whole-cell bacterial bioreporter sensors for aromatic compounds are being developed. In a bioreporter sensor, a genetically encoded reporter protein is produced in response to a contact of a microbial cell with an analyte [25–27]. The main limiting factor for construction of whole-cell bacterial bioreporter sensors for chloroaromatic compound is shortage of knowledge about metabolic (biochemical) pathways of bacteria capable of degrading chloro-substituted aromatic compounds [26].

In this study, laboratory models of the membrane microbial sensor were formed on the basis of immobilized cells of *R. opacus* ICP, which contained enzymes of degradation of benzoate or 3-chlorobenzoate (benzoate 1,2-dioxygenase or 3-chlorobenzoate 1,2-dioxygenase, respectively), to explore features of microbial metabolism of toxic aromatic compounds. The models were applied to evaluate the actinobacterial cells' responses to benzoate and 3-chlorobenzoate in the absence of both enzymes, benzoate 1,2-dioxygenase and 3-chlorobenzoate 1,2-dioxygenase, or in the presence of one of the said enzymes.

## 2. Materials and methods

### 2.1 The microorganism

The object of our study was a Gram-positive non-spore-forming *R. opacus* ICP actinobacterium (DSM 46757, and VKM Ac-2638), which was isolated from the selective medium with 2,4-dichlorophenol. The culture was able to degrade benzoate (BA) and some of substituted benzoates [15]. The actinobacterium was maintained on agarized Luria-Bertani (LB) medium and transferred every 6 months.

## 2.2 Culture conditions

The culture was grown on Petri dishes containing agarized LB medium (the medium without 3CBA and BA) or agarized mineral medium with BA (200 mg/L) as a sole carbon and energy source. The mineral medium had the following composition (g/L):  $\text{Na}_2\text{HPO}_4$  – 0.73;  $\text{KH}_2\text{PO}_4$  – 0.35;  $\text{MgSO}_4 \times 7\text{H}_2\text{O}$  – 0.1;  $\text{NaHCO}_3$  – 0.25;  $\text{MnSO}_4$  – 0.002;  $\text{NH}_4\text{NO}_3$  – 0.75;  $\text{FeSO}_4 \times 7\text{H}_2\text{O}$  – 0.02. Petri dishes were incubated at 28 °C until bacterial cells germination. Biomass grown was washed out with liquid LB or mineral medium, depending on the growth medium, and cell suspensions prepared were used as inoculums. To produce biomass, inoculum suspension was inoculated in 750-mL Erlenmeyer flasks with 200 mL of appropriate liquid growth medium: liquid LB medium (for LB-grown cells, which were BDO- and 3CBDO-free cells) or liquid mineral medium with 200 mg/L of BA (for BA-grown cells). The culture was grown at 28 °C on a rotary shaker ( $n = 220$  rpm).

Grown *R. opacus* 1CP cells were centrifuged (12,000–16,000 g, 10–15 min, + 4 °C) and washed twice with a 50-mM K-Na phosphate buffer (pH 7.4). BA-grown biomass and one part of LB-grown biomass were suspended in the phosphate buffer up to 100 mg of wet cells per mL, stored at + 4 °C for 12 h and were then used for preparation of receptor elements on the basis of BA- or LB-grown cells, respectively. Another part of LB-grown biomass was induced by BA or 3CBA.

## 2.3 BDO or 3CBDO induction in LB-grown *R. opacus* 1CP cells

To induce BDO or 3CBDO in LB-grown *R. opacus* 1CP cells, LB-grown biomass was suspended in the phosphate buffer containing 2 g/L of BA (for BA-induced cells) or in the phosphate buffer containing 100 mg/L of 3CBA (for 3CBA-induced cells), respectively. The culture cells were incubated at 28 °C on a rotary shaker ( $n = 220$  rpm) for 96 h (for BA-induced cells) or 24 h (for 3CBA-induced cells). After incubation, biomass (BA- or 3CBA-induced biomasses) was centrifuged and washed twice with the phosphate buffer. BA- and 3CBA-induced biomass were suspended in the phosphate buffer up to 100 mg of wet cells per mL, stored at + 4 °C for 12 h and were then used for preparation of receptor elements based on BA- or 3CBA-induced cells, respectively.

## 2.4 Formation of bioreceptors based on *R. opacus* 1CP cells

Bioreceptors were formed on the basis of obtained suspensions (100 mg of wet cells weight per mL) of the actinobacterial cells. To form a bioreceptor, bacterial suspension was immobilized on Whatman paper by the method of physical adsorption. To achieve it, actinobacterial suspension (10  $\mu\text{L}$ ) was spotted (3–4 mm in diameter) onto a piece of paper (4  $\times$  4 mm<sup>2</sup>). Prepared bioreceptors were air-dried for 30–40 min and used for formation of laboratory models

of the biosensor or stored at + 4 °C.

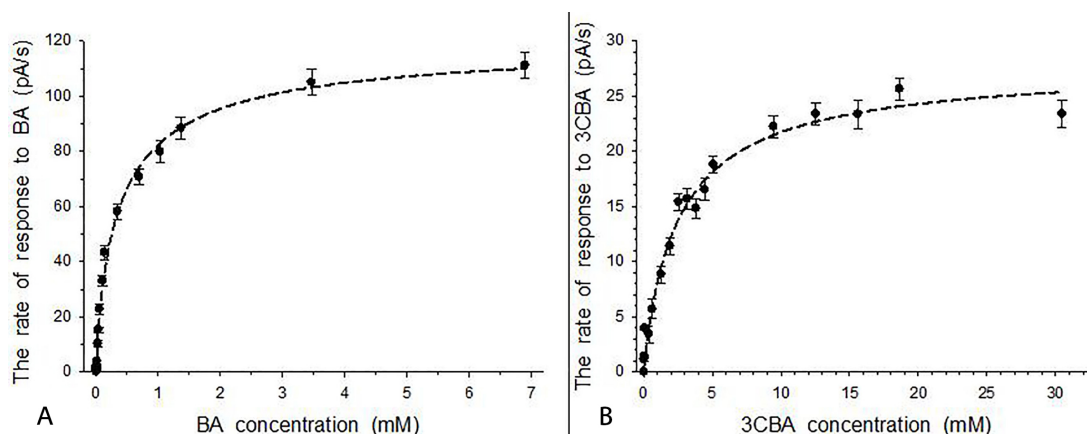
## 2.5 Formation of laboratory models of the membrane microbial sensor

Cells' responses to substrates were determined by biosensor method using laboratory models of the membrane microbial sensor. Each laboratory model was formed on the basis of the bioreceptor using *R. opacus* cells either LB- or BA-grown, BA- or 3CBA-induced cells. To form a recognizing part of the biosensor, the bioreceptor was fixed on the measuring surface of the Clark-type oxygen electrode by means of nylon net. The oxygen electrode with the bioreceptor (a microbial electrode) was placed in an open 5 mL measuring cell. The cell was filled with an air-saturated buffer and equipped with a magnetic stirrer. The design of the recognizing part of the membrane microbial sensor is presented in our previously published paper [28]. The Clark-type oxygen electrode was used as a transducer of the response of microbial cells. An amplifier system (Ingold 531-04 O<sub>2</sub> Amplifier, Switzerland-USA) amplified the signal of the oxygen electrode. Cells' response to substrate was registered with a two-coordinate recorder (XY Recorder-4103, Czech Republic).

## 2.6 Determination of a response to substrate (BA or 3CBA) for immobilized *R. opacus* 1CP cells used as the recognizing element of the membrane microbial sensor

The measurements of cells' responses to substrates were performed in the air-saturated phosphate buffer at + 20–22 °C at permanent stirring. The microbial electrode was placed in the measuring cell with the buffer solution and basal (endogenous) cell respiration was then stabilized. After stabilization of endogenous respiration of *R. opacus* cells, substrate solution was injected into the buffer solution. The cells' response to substrate was proportional to a rate of cells' respiration change. The change of cells' respiration led to a change in oxygen concentration at the measuring area of the Clark-type oxygen electrode. The Clark-type oxygen electrode transduced a chemical signal (the change in oxygen concentration) into electric signal (a change in the electrode current). The change in the electrode current was recorded with the two-coordinate recorder. The recorded signal reflected the rate of change in the electrode current ( $dI/dt$ , pA/s), which was proportional to a change in oxygen consumption by culture in response to substrate injection. The cells' response to substrate was calculated as the first derivative of the electrode current change in response to substrate addition (pA/s). After measurement, the system was washed, the basal respiration was then registered, and the system was ready to make the next measurement.

To measure the response to substrate for another *R. opacus* 1CP cells using another bioreceptor, an appropriate bioreceptor was fixed on the measuring surface of the Clark-type oxygen electrode, and measurements were made.



**Fig. 2. Responses to the substrate for immobilized *R. opacus* 1CP cells grown in LB medium (BA- and 3CBA-free).** (A) Responses to BA for BDO-free cells. (B) Responses to 3CBA for 3CBDO-free cells. Error bars indicate standard deviation from the measurement of three samples.

## 2.7 Statistics

The presented data are the results of one of two/three independent experiments. The measurements were taken in triplicate. Presented results are average values. Statistical data analysis was carried out using a Student's *t*-test taking  $p < 0.05$ .

## 3. Results and discussion

### 3.1 LB-grown (non-induced by BA or 3CBA) cells' responses to substrates for characterization of BA and 3CBA transport into cells

Our earlier study showed that BDO and 3CBDO of *R. opacus* 1CP are inducible enzymes, which are synthesized in the cells grown in the presence of substrates of enzymes [23,29]. Furthermore, in our previous work, the response to the substrate for intact *R. opacus* 1CP cells used as a recognizing element of a reactor microbial sensor was an indicator of the activity of the enzyme initiating degradation of the substrate, BDO or 3CBDO [24]. LB medium is the BA- and 3CBA-free medium. Therefore, LB-grown actinobacterial cells contained only insignificant amounts of inducible enzymes (basal activity), which could not be registered in microbial cells.

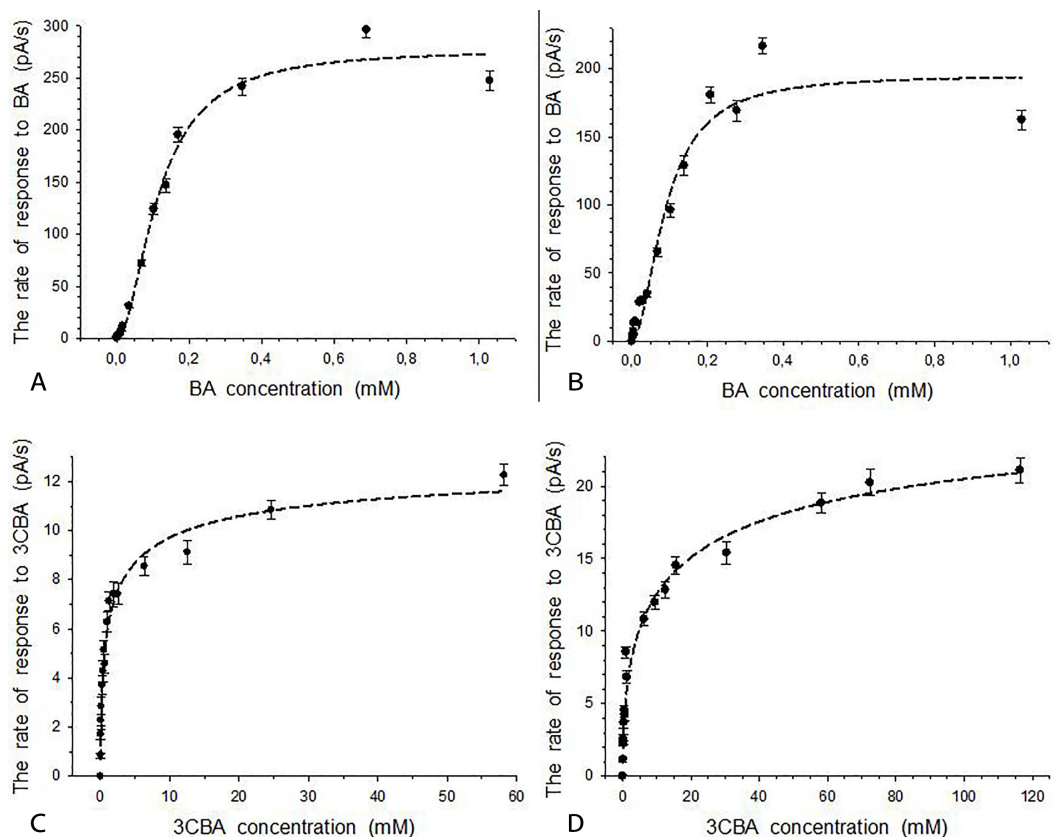
It is known that the response to a substrate for immobilized cells of a bioreceptor of a membrane microbial sensor is caused by processes of both transport of a substrate into microbial cells and metabolism of a substrate in cells [30]. It is obvious that in the absence of the enzyme, which initiates substrate degradation, immobilized cells' response to a substrate is caused by processes of substrate transport into cells. When no BDO and 3CBDO activity were registered in intact LB-grown *R. opacus* 1CP cells, immobilized cells' response to BA or 3CBA was caused by transport of a substrate into LB-grown cells. Thus, curves shown in Fig. 2 describe a change in the rate of BA and 3CBA transport into *R. opacus* 1CP cells. Detection of the response to BA

and 3CBA for 1CP actinobacterial cells non-induced by the substrate was the evidence of the presence of a constitutive system for substrates transport into cells. Earlier, both for Gram-negative bacteria and substrate other than BA, permeability of non-induced and induced cells to substrate was registered by Hegeman for *Pseudomonas putida* ATCC 12633/mandelate [31] as well as a constitutive system of uptake was observed by Miguez *et al.* [32] for *Alcaligenes denitrificans* BRI 6011/2,4-dichlorobenzoic acid.

LB-grown cells of the culture are BDO- and 3CBDO-free cells. When BA and 3CBA penetrate the LB-grown cells, these substrates cannot be metabolized by the cells. However, these substrates were not accumulated in the cells. It was supported by the fact that receptors based on immobilized *R. opacus* 1CP cells, which was not induced by substrate, operated for longer than 24 h without loss of activity and sample throughput [33] of the membrane microbial sensor was 4–5 samples per hour. Hence, BA and 3CBA should have entered and exited LB-grown 1CP cells. As for substrate-induced cells of *Pseudomonas putida* ATCC 12633 in the presence of mandelate, concentrating permease activity was shown by Higgins and Mandelstam [34]. Since BA and 3CBA were not accumulated in the cells, constitutive concentrating permease could not mediate transport of these substrates into the cells of 1CP culture.

Aubert and Motais demonstrated that weak benzoic acid penetrates the cell through the membrane lipids as undissociated acid [35]. In the present study, cells' responses were measured in the buffer at pH 7.4. In aqueous solution at pH 7.0, BA was dissociated by approximately 99.9% [36], and significant amounts of BA (as anion) were unable to independently diffuse into the cell through simple diffusion. Saturation kinetics (Fig. 2) was observed for dependency of the rates of *R. opacus* 1CP response to BA or 3CBA on BA or 3CBA concentration, respectively. It





**Fig. 3. Responses to BA or 3CBA for immobilized *R. opacus* 1CP cells in the presence of BDO in cells.** (A) Responses to BA for BA-induced cells. (B) Responses to BA for BA-grown cells. (C) Responses to 3CBA for BA-induced cells. (D) Responses to 3CBA for BA-grown cells. Error bars indicate standard deviation from the measurement of three samples.

excluded the involvement of simple (passive) diffusion in BA and 3CBA transport into LB-grown 1CP cells (linear regression between the rate and concentration would be obtained). At the same time, active transport of BA and 3CBA was hardly improbable for resting immobilized cells knowing that the active transport is coupled with ATP hydrolysis. Most likely, carriers-mediated diffusion facilitated BA and 3CBA transport into non-induced 1CP cells. The question remains unanswered whether porins like BenF and BenP (porins can also be present in a wall zone of Gram-positive bacteria [37]) and membrane BA transport proteins, such as BenE1 and BenE2 (2.A.46 benzoate:H<sup>+</sup> symporter BenE family) [38–40], could participate in the transport of BA and 3CBA into 1CP cells non-induced by the substrate.

Nonlinear regression fit of data to Michaelis-Menten equation resulted in calculation of rate constants  $V_{max}$  of  $109.0 \pm 2.7$  pA/s and  $27.2 \pm 0.9$  pA/s and transport affinity constants  $S_{0.5}$  (as  $K_m$ ) of  $0.27 \pm 0.03$  mM and  $2.44 \pm 0.29$  mM for BA and 3CBA, respectively. Two explanations can be given for the results obtained. First, these values could indicate not narrow specificity of the single constitutive system mediating substrate (both BA and 3CBA) transport into cells of 1CP actinobacterium. The affinity of the transport system was higher to BA (0.27 mM) than

to 3CBA (2.44 mM). The rate of BA transport (Fig. 2A) into resting *R. opacus* 1CP cells was higher than the rate of 3CBA transport (Fig. 2B). Second, probably two different transport systems are involved in transport of BA or 3CBA into non-induced 1CP cells. For instance, Chaudhry *et al.* [38] reported that the rates of BA uptake into resting *Corynebacterium glutamicum* cells were higher when BenE, but not BenK, mediated BA transport. In addition, both specific and non-specific pore-forming proteins have been found in Gram-positive bacteria [41].

### 3.2 The responses to substrates for BA-induced and BA-grown cells

Both cells induced with BA in non-growth conditions (referred as BA-induced cells) and cells grown in BA-medium (referred as BA-grown cells) contained BDO initiating BA metabolism in cells. Therefore, the response to substrate for BDO-containing immobilized *R. opacus* 1CP cells was caused by both the process of substrate transport into cells and process of initial metabolism of substrate in cells. Due to specificity of BDO of 1CP actinobacterium, responses to both substrates (BA and 3CBA), which characterized BDO activity, were registered earlier using intact cells [10]. In the present study, responses to BA (Fig. 3A and 3B) and 3CBA (Fig. 3C and 3D) were studied us-

ing BDO-containing immobilized cells of the culture. The cells' response to BA was markedly higher than the cells' response to 3CBA. It could be a result of a drop of enzyme activity (BDO activity) in the presence of 3CBA as a substrate. Similar change of enzyme activity was shown in previous research with intact cells: BDO activity in the presence of BA was higher by a factor of 5 than in the presence of 3CBA [10]. It is a similar situation as described by Krooneman *et al.* [42] for determination of maximum oxygen uptake rate in the presence of BA or 3CBA for BA-grown *Alcaligenes* sp. L6. For this culture in the presence of BA, cells' response was a 5-fold higher than a response in the presence of 3CBA. The difference between cells' responses to BA and 3CBA can also be explained by the presence of different transporters in ICP cells in the presence of BA or CBA. For instance, various transporters were induced by BA (BenK, BenE1 and BenE2) and 3CBA (BenK and BenE2) in *Pseudomonas putida* KT2440 cells [43]. Furthermore, in confirmation of the above said about the response to substrate for immobilized ICP cells, the categories of expressed genes such as 'energy metabolism' and 'transport and binding proteins' characterized the cellular responses to substrate for *Pseudomonas putida* KT2440 cells reported by Wang *et al.* [43]. In addition, Clark *et al.* and Wong *et al.* [40,44] showed that transport process and process of substrate metabolism are dependent of each other.

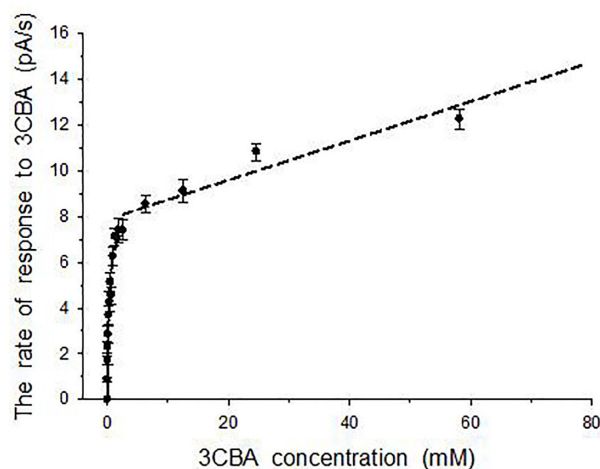
Curves of dependencies of the cells' response to substrate (both BA and 3CBA) on substrate concentration for BA-induced and BA-grown immobilized cells are represented in Fig. 3. Saturation curves were obtained. Non-linear regression fit of data to the Hill equation was found for all curves. Half-saturation constants,  $S_{0.5}$ , which were integrated constants of affinity to substrate for processes causing immobilized cells' response to substrate, were calculated. For BDO of *R. opacus* ICP cells, enzyme specificity to BA was higher than specificity to 3CBA [10]. As expected, in the presence of BDO for both BA-induced and BA-grown cells, constants of affinity to BA were lower than the constants to 3CBA, affinity to BA was higher than that to 3CBA. So, for BA-induced cells, the constant of affinity to BA ( $117 \mu\text{M}$ ) was by an order lower than that to 3CBA ( $1400 \mu\text{M}$ ). For BA-grown cells, affinity to BA virtually did not change, namely the constant of affinity to BA was insignificantly lower ( $91 \mu\text{M}$ ) than that for BA-induced cells.

Regarding the Hill kinetics for cells' responses to BA, positive kinetic cooperativity by the substrate was found for responses of BA-induced and BA-grown immobilized cells. Values of Hill coefficient ( $n$ ) were greater than 1. They were 1.87 and 1.90 for BA-induced and BA-grown ICP cells, respectively. It was in conformity with dependency of cooperatives by the substrate for BDO on BA concentration in the growth medium for *R. opacus* ICP cells. This phenomenon was found in our study using intact cells [45]. For BDO of intact ICP cells, positive kinetic cooper-

ativity by BA was detected when BA concentration in the growth medium was lower than 6 g/L. Furthermore, considering the dependency between substrate transport process and process of substrate metabolism [44], positive kinetic cooperativity was observed by Choudhary *et al.* [38] for BA transport into *Pseudomonas putida* CSV86 cells ( $n = 1.9$ ).

For cells' responses to 3CBA in the presence of BDO, unlike cells' responses to BA, negative kinetic cooperativities by the substrate were detected.  $n$  values were 0.54 and 0.51 for BA-induced and BA-grown ICP cells, respectively. Responses to 3CBA were measured for the same cells induced by BA, which were used for detection of responses to BA. In both experiments (both with BA and with 3CBA) the same enzyme (BDO) caused the cells' response to substrate. Therefore, allosteric interaction of active sites of BDO in the presence of 3CBA could be a reason of negative kinetic cooperativities by substrate [46].

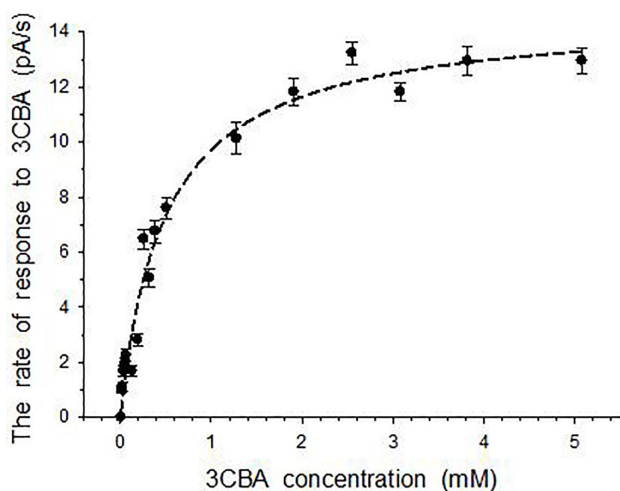
Another reason of negative kinetic cooperativity could be the presence of two phases (hyperbolic and linear) for 'response-3CBA concentration' dependency that was obtained for BA-induced cells (Fig. 4). Linear phase indicated that the simple diffusion took part in formation of the cells' response to 3CBA when 3CBA concentration was higher than 10 mM.



**Fig. 4.** Two phases dependency of the rate of the cells' response to 3CBA on 3CBA concentration for BA-induced *R. opacus* ICP cells. Error bars indicate standard deviation from the measurement of three samples.

### 3.3 The responses to 3CBA for 3CBA-induced cells

After cells induction in the presence of 3CBA, *R. opacus* ICP cells contained 3CBDO. For these immobilized cells, the response to 3CBA was caused by 3CBA transport and 3CBDO activity. A curve of the dependency of the cells' response to 3CBA on concentration of 3CBA is shown in Fig. 5.



**Fig. 5. Responses to 3CBA for immobilized *R. opacus* 1CP cells induced by 3CBA.** Error bars indicate standard deviation from the measurement of three samples.

Saturation curve was obtained; data of nonlinear regression fitted to the Michaelis-Menten equation. Half-saturation constants ( $S_{0.5}$ ), which characterized the process affinity to 3CBA, were calculated using the Michaelis-Menten model. 3CBA substrate affinity to 'its' enzyme, 3CBDO, should be higher than that to BDO. For instance, enzyme activity of 3CBA-grown *Alcaligenes* sp. L6 cells was higher in the presence of 3CBA but not in the presence of BA [42]. Hence, integrated constant of process affinity to 3CBA should be lower for 3CBA-induced 1CP cells in comparison with BA-induced the cells. That is what we observed for 3CBA-induced *R. opacus* 1CP cells. For a 3CBA-3CBDO combination, calculated half-saturation constant ( $S_{0.5}$ , 484  $\mu$ M) was the same order as the value determined for a BA-BDO combination (117  $\mu$ M) and was on order of magnitude less than for 3CBA-BDO.

#### 4. Conclusions

In sum, laboratory models of the membrane microbial sensor, based on immobilized *Rhodococcus opacus* 1CP cells and the Clark-type oxygen electrode as a transducer, were successfully applied to evaluate the actinobacterial cells' responses to BA and 3CBA.

The data from this study show that the membrane microbial sensor based on immobilized cells of the bacterial culture could be useful as a perspective tool for evaluation of enzymes of complex structure initiating substrate metabolism coupled with oxygen consumption.

The *Rhodococcus opacus* 1CP culture has constitutive system(s) for the transport of BA and 3CBA into cells. Therefore, cells without prior induction can be introduced into an environment contaminated with these xenobiotics. The positive cooperativity by the substrate, which was shown for immobilized cells of the culture, indicates that the catalytic efficiency of the active sites of the enzyme

(BDO) increases as they are filled with the substrate. Moreover, adaptation to one substrate (BA or 3CBA) does not preclude the use of the second substrate. In the light of it, we can conclude that *Rhodococcus opacus* 1CP is an ideal xenobiotics-degrading culture for remediation of soil contaminated by BA and 3CBA.

The results of the study have indicated a direction for our further research. Previously, the presence of porins was reported only for Gram-negative bacteria. Currently, pore-forming proteins have been found in Gram-positive bacteria, including rhodococci. There is no data on the pore-forming proteins of *Rhodococcus opacus* 1CP. Microbiological studies are planned to confirm or refute the presence of pore-forming proteins in the wall zone of Gram-positive *Rhodococcus opacus* 1CP. In addition, genetic studies will help identify genes encoding BA and 3CBA transporters.

#### Abbreviations

BA, benzoate; BDO, benzoate 1;2-dioxygenase; 3CBA, 3-chlorobenzoate; 3CBDO, 3-chlorobenzoate 1;2-dioxygenase.

#### Author contributions

EVE and IPS conceived and designed the experiments, and performed the experiments; EVE analyzed the data; EVE and IPS contributed reagents and materials. Both authors read and approved the final manuscript.

#### Ethics approval and consent to participate

Not applicable.

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#### Conflict of interest

The authors declare no conflict of interest.

#### References

- [1] Molnari JC, Myers AL. Carbonyl reduction of bupropion in human liver. *Xenobiotica*. 2012; 42: 550–561.
- [2] Field JA, Sierra-Alvarez R. Microbial transformation of chlorinated benzoates. *Reviews in Environmental Science and Bio/Technology*. 2008; 7: 191–210.
- [3] Kamei I, Kogura R, Kondo R. Metabolism of 4,4'-dichlorobiphenyl by white-rot fungi *Phanerochaete chrysosporium* and *Phanerochaete* sp. MZ142. *Applied Microbiology and Biotechnology*. 2006; 72: 566–575.
- [4] Pieper DH. Aerobic degradation of polychlorinated biphenyls. *Applied Microbiology and Biotechnology*. 2005; 67: 170–191.
- [5] Niedan V, Schöler HF. Natural formation of chlorobenzoic acids (CBA) and distinction between PCB-degraded CBA. *Chemosphere*. 1997; 35: 1233–1241.

- [6] Kasberg T, Daubaras DL, Chakrabarty AM, Kinzelt D, Reineke W. Evidence that operons *tcb*, *ifd*, and *clc* encode maleylacetate reductase, the fourth enzyme of the modified ortho pathway. *Journal of Bacteriology*. 1995; 177: 3885–3889.
- [7] Commandeur LC, Parsons JR. Degradation of halogenated aromatic compounds. *Biodegradation*. 1990; 1: 207–220.
- [8] Reineke W, Knackmuss HJ. Microbial degradation of haloaromatics. *Annual Review of Microbiology*. 1988; 42: 263–287.
- [9] Hartmann J, Reineke W, Knackmuss HJ. Metabolism of 3-chloro-, 4-chloro-, and 3,5-dichlorobenzoate by a pseudomonad. *Applied and Environmental Microbiology*. 1979; 37: 421–428.
- [10] Solyanikova IP, Emelyanova EV, Shumkova ES, Egorova DO, Korsakova ES, Plotnikova EG, *et al.* Peculiarities of the degradation of benzoate and its chloro- and hydroxy-substituted analogs by actinobacteria. *International Biodeterioration & Biodegradation*. 2015; 100: 155–164.
- [11] Čejková A, Masák J, Jirků V, Veselý M, Pátek M, Nešvera J. Potential of *Rhodococcus erythropolis* as a bioremediation organism. *World Journal of Microbiology and Biotechnology*. 2005; 21: 317–321.
- [12] Ajithkumar PV, Kunhi AA. Pathways for 3-chloro- and 4-chlorobenzoate degradation in *Pseudomonas aeruginosa* 3mT. *Biodegradation*. 2000; 11: 247–261.
- [13] Hernandez BS, Higson FK, Kondrat R, Focht DD. Metabolism of and inhibition by chlorobenzoates in *Pseudomonas putida* P111. *Applied and Environmental Microbiology*. 1991; 57: 3361–3366.
- [14] Solyanikova IP, Emelyanova EV, Shumkova ES, Travkin VM. Pathways of 3-chlorobenzoate degradation by *Rhodococcus opacus* strains ICP and 6a. *Microbiology*. 2019; 88: 563–572.
- [15] Gorlatov SN, Maltseva OV, Shevchenko VI, Golovleva LA. Degradation of chlorophenols by a culture of *Rhodococcus erythropolis*. *Microbiology*. 1989; 58: 647–651.
- [16] Dorn E, Hellwig M, Reineke W, Knackmuss HJ. Isolation and characterization of a 3-chlorobenzoate degrading pseudomonad. *Archives of Microbiology*. 1974; 99: 61–70.
- [17] Nakatsu CH, Straus NA, Wyndham RC. The nucleotide sequence of the Tn5271 3-chlorobenzoate 3,4-dioxygenase genes (*cbaAB*) unites the class IA oxygenases in a single lineage. *Microbiology*. 1995; 141: 485–495.
- [18] Romanov V, Hausinger RP. *Pseudomonas aeruginosa* 142 uses a three-component ortho-halobenzoate 1,2-dioxygenase for metabolism of 2,4-dichloro- and 2-chlorobenzoate. *Journal of Bacteriology*. 1994; 176: 3368–3374.
- [19] Huang D, Whang T, Cheng F, Wu Y, Wang Y, Luo W, *et al.* Toxicity assessment of mono-substituted benzenes and phenols using a *Pseudomonas* initial oxygen uptake assay. *Environmental Toxicology and Chemistry*. 2005; 24: 253–260.
- [20] Hickey WJ, Focht DD. Degradation of mono-, di-, and trihalogenated benzoic acids by *Pseudomonas aeruginosa* JB2. *Applied and Environmental Microbiology*. 1990; 56: 3842–3850.
- [21] Farr DR, Cain RB. Catechol oxygenase induction in *Pseudomonas aeruginosa*. *The Biochemical Journal*. 1968; 106: 879–885.
- [22] Tizzard AC, Bergsma JH, Lloyd-Jones G. A resazurin-based biosensor for organic pollutants. *Biosensors & Bioelectronics*. 2006; 22: 759–763.
- [23] Emelyanova EV, Solyanikova IP. Evaluation of 3-chlorobenzoate 1,2-dioxygenase inhibition by 2- and 4-chlorobenzoate with a cell-based technique. *Biosensors-Basel*. 2019; 9: 106.
- [24] Solyanikova IP, Borzova OV, Emelyanova EV. Kinetics of interaction between substrates/substrate analogs and benzoate 1,2-dioxygenase from benzoate-degrading *Rhodococcus opacus* ICP. *Folia Microbiologica*. 2017; 62: 355–362.
- [25] Alhadrami HA. Biosensors: Classifications, medical applications, and future prospective. *Biotechnology and Applied Biochemistry*. 2018; 65: 497–508.
- [26] Plotnikova EG, Shumkova ES, Shumkov MS. Whole-cell bacterial biosensors for the detection of aromatic hydrocarbons and their chlorinated derivatives. *Prikladnaia Biokhimiia i Mikrobiologiya*. 2018; 52: 347–357.
- [27] Close DM, Ripp S, Saylor GS. Reporter proteins in whole-cell optical bioreporter detection systems, biosensor integrations, and biosensing applications. *Sensors*. 2009; 9: 9147–9174.
- [28] Emelyanova EV, Solyanikova IP. Evaluation of phenol-degradation activity of *Rhodococcus opacus* ICP using immobilized and intact cells. *International Journal of Environmental Science and Technology*. 2020; 17: 2279–2294.
- [29] Solyanikova IP, Borzova OV, Emelyanova EV, Shumkova ES, Prisyazhnaya NV, Plotnikova EG, *et al.* Dioxygenases of chlorobiphenyl-degrading species *Rhodococcus wratislaviensis* G10 and chlorophenol-degrading species *Rhodococcus opacus* ICP induced in benzoate-grown cells and genes potentially involved in these processes. *Biochemistry*. 2016; 81: 986–998.
- [30] Turner APF, Karube I, Wilson GS. *Biosensors: fundamentals and applications*. Oxford University Press: New York. 1987.
- [31] Hegeman GD. Synthesis of the enzymes of the mandelate pathway by *Pseudomonas putida*. *Journal of Bacteriology*. 1966; 91: 1155–1160.
- [32] Miguez CB, Greer CW, Ingram JM, Macleod RA. Uptake of benzoic acid and chloro-substituted benzoic acids by *Alcaligenes denitrificans* BRI 3010 and BRI 6011. *Applied and Environmental Microbiology*. 1995; 61: 4152–4159.
- [33] Thévenot DR, Toth K, Durst RA, Wilson GS. Electrochemical biosensors: recommended definitions and classification. *Biosensors & Bioelectronics*. 2001; 16: 121–131.
- [34] Higgins SJ, Mandelstam J. Evidence for induced synthesis of an active transport factor for mandelate in *Pseudomonas putida*. *The Biochemical Journal*. 1972; 126: 917–922.
- [35] Aubert L, Motais R. Molecular features of organic anion permeability in ox red blood cell. *The Journal of Physiology*. 1975; 246: 159–179.
- [36] Holyoak CD, Bracey D, Piper PW, Kuchler K, Coote PJ. The *Saccharomyces cerevisiae* weak-acid-inducible ABC transports fluorescein and preservative anions from the cytosol by an energy-dependent mechanism. *Journal of Bacteriology*. 1999; 181: 4644–4652.
- [37] Costa-Riu N, Burkovski A, Krämer R, Benz R. PorA represents the major cell wall channel of the Gram-positive bacterium *Corynebacterium glutamicum*. *Journal of Bacteriology*. 2003; 185: 4779–4786.
- [38] Choudhary A, Purohit H, Phale PS. Benzoate transport in *Pseudomonas putida* CSV86. *FEMS Microbiology Letters*. 2017; 364.
- [39] Chaudhry MT, Huang Y, Shen X, Poetsch A, Jiang C, Liu S. Genome-wide investigation of aromatic acid transporters in *Corynebacterium glutamicum*. *Microbiology*. 2007; 153: 857–865.
- [40] Clark TJ, Momany C, Neidle EL. The benPK operon, proposed to play a role in transport, is part of a regulon for benzoate catabolism in *Acinetobacter* sp. strain ADP1. *Microbiology*. 2002; 148: 1213–1223.
- [41] Achouak W, Heulin T, Pagès JM. Multiple facets of bacterial porins. *FEMS Microbiology Letters*. 2001; 199: 1–7.
- [42] Krooneman J, Wieringa EB, Moore ER, Gerritse J, Prins RA, Gottschal JC. Isolation of *Alcaligenes* sp. strain L6 at low oxygen concentrations and degradation of 3-chlorobenzoate via a pathway not involving (chloro)catechols. *Applied and Environmental Microbiology*. 1996; 62: 2427–2434.
- [43] Wang Y, Morimoto S, Ogata N, Fujii T. A survey of cellular responses in *Pseudomonas putida* KT2440 growing in sterilized soil by microarray analysis. *FEMS Microbiology Ecology*. 2011; 78: 220–232.
- [44] Wong CM, Dilworth MJ, Glenn AR. Cloning and sequencing show that 4-hydroxybenzoate hydroxylase (PobA) is required for uptake of 4-hydroxybenzoate in *Rhizobium leguminosarum*. *Microbiology*. 1994; 140: 2775–2786.
- [45] Emelyanova EV, Solyanikova IP. Benzoate concentration and cooperativity by a substrate for benzoate 1,2-dioxygenase from benzoate-degrading *Rhodococcus opacus* ICP. *Journal of Biotechnology and Biomedical Science*. 2017; 1: 38–46.
- [46] Kurganov BI. *Allosteric enzymes: Kinetic behavior*. Wiley: New York. 1983.