#### иилогонхадона виньодения и киномительной импорительной виньодения виньодения

# Surfactant degradative plasmids

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It has been shown that the degradation of anionic, cationic and ampholytic surfactants by pseudomonads can be controlled by the plasmids with size of 60—130 kb. Most plasmid strains that degrade surfactants are capable of conjugative transfer and elimination from bacteria cells. Restriction patterns of plasmids do not reveal significant homology between plasmids.

Introduction. The data about microorganisms able to utilize cationic and anionic surfactants for metabolism are insufficient. The genetic control of their degradation has not been clearly determined yet.

The plasmid localization of the genes responsible for degradation of an anionic surfactant, alkyl benzenesulfonate (ABS), was first shown by Caine et al. [1]. They discovered a conjugative plasmid controlling at least one step of ABS degradation—the cleavage of sulfonic group in the strain-degrader Pseudomonas testosteroni. Though the biochemical aspects of this problem had been investigated [2], further studies on the genetic control and regulation of the ABS degradation were not made.

This is the only report of a possible plasmid control in the case of ABS and of chromosomal localization of degradative genes for naphthalene sulfonates [3].

Studies on the localization of genetic determinants controlling surfactant degradation expand our knowledge about microorganism-degraders. The data obtained may be used for construction of new strains-destructors which could be perspective for the waste water treatment. This is important for the development of biotechnologies aimed at purifying waste water from these toxicants.

We have selected and characterized highly active bacterial strains which degrade ionogenic surfactants

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referring to different classes [4, 5]. Some of them have been tested for the ability to purify waste water on the industrial scale [6, 7].

The localization of the genetic determinants controlling transformation of ionogenic surfactants by bacteria-degraders has been shown.

Materials and Methods. The following surfactants were used in the study:

Anionic surfactants

Volgonat,  $[(C_nH_{2n+1}C_mH_{2m+1})CHSO_3]$ , containing 40 % of the major compound (sodium alkansulfonate), where n+m=11-17. The sample of volgonat contains nonsulfonated compounds, NaCl (6 %), Na<sub>2</sub>SO<sub>4</sub> (1 %), Na<sub>2</sub>SO<sub>3</sub> (0.006 %), Fe (0.8 %), NaOH (0.1 %).

Alkylsulfate,  $\{C_nH_{2n+1}OSO_3Na\}$ , containing 30 % of the major compound, where n = 16-18,  $Na_2SO_4$  (2 %).

DNS-A, [NaO<sub>3</sub>SCH(COONa)CH<sub>2</sub>COOCH<sub>2</sub>CH<sub>2</sub>-N(H)OOC<sub>n</sub>H<sub>2n+1</sub>], containing 35 % of the major compound (disodium mono-2-(alkanoamino)ethylsuccinatosulfonate, where n = 9-12. The sample contains nonsulfonated compounds, Na<sub>2</sub>SO<sub>4</sub> (7 %) and Na<sub>2</sub>SO<sub>3</sub> (2 %)

Metaupon,  $[C_nH_{2n+1}CONHC_2H_4SO_3Na]$ , containing 50 % of the major compound.

ABS(chloric sulfonol),  $[C_nH_{2n+1}C_6H_4SO_3Na]$ , containing 40 % of the major compound (alkylbenzenesulfonate) obtained on the kerosene basis, where n = 10-13. Its fractional composition:  $C_{1-10} - 20$  %,  $C_{11-18} - 80$  %. The sample of ABS contains nonsul-

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fonated compounds (2 %),  $Na_2SO_4$  (6 %) and Fe (0.006 %).

SDS (sodium dodecylsulphate),  $[C_{11}H_{23}CHSO_3-Na]$ , containing 99 % of the major compound. Ampholytic surfactants

Cyclimide, containing the major compound (30 %), where NaCl (10—11 %) and moisture (nearly 60 %).

Amidobethaine (trisodium N,N,N'-triacetate-2-[N-acyl-N-(2-hydroxyethyl)-amino lethyl ammonium hydroxide), containing the major compound (40 %), where NaCl (14.5 %) and moisture (45 %).

$$\begin{bmatrix} HOCH_2CH_2 \\ C_nH_{2n+1}CO \end{bmatrix} NCH_2CH_2N^{\dagger}(CH_2COONa)_3$$
 OH

Alkyl aminobispropionate (disodium N-alkylaminobis (3,3'-propionate) (AABP), containing the major compound (85-88%), where n=18-20, amines (1%).

Cationic surfactants

Catamine AB { $[C_nH_{2n+1}N^+(CH_3)_2CH_2C_6H_5]C\Gamma$ }, containing the major compound (50 %), where n = 10-18, tertiary amines (0.6 %), salts of tertiary amines (1.8 %, moisture (nearly 47 %).

Alkamone DC {  $[C_nH_{2n+1}OCH_2N^+(CH_3)(CH_2-CH_3)_2][CH_3SO_4^-]$ , containing the major compound (90 %), where n = 10-18, secondary amines (2.7—3.5 %), moisture (nearly 6.5 %).

Ethonium, (1,2-[N,N-bis(dimethyl)-N,N'-bis(decylacetate)ethylene diammonium dichloride).

$$\begin{array}{c|c} (CH_3)_2N^{\dagger}(CH_2)_2N^{\dagger}(CH_3)_2 & \textbf{2CI} \\ & & & \\ CH_{10}H_{21}COOCH_2 & CH_2COOCH_{10}H_{21} \\ \end{array}$$

The bacterial strains-degraders of surfactants are given in Table. Bacteria were grown in LB broth and agar. The medium of the following composition  $(g \cdot l^{-1})$  was used as minimal:  $K_2HPO_4-1.0$ ;  $NH_4NO_3-1.0$ ; KCl-0.5;  $MgCl_2-0.1$ . 15 g of agar was added to obtain agarized medium. Anionic and cationic surfactants  $(0.3 \ g \cdot l^{-1})$  were employed as the sole carbon and energy source.

Strain Escherichia coli K802 [14] were grown at 37 °C, other strains at 28 °C.

In the work the following plasmids have been employed: pBS222 A-15 (Tc', Cm'), pLAFR5 (21,5 kb) [15], pRK2013 (Rep ColE1) [16], RP4 (55 kb) [17], NPL-1 (98 kb), pBS2 (130 kb).

To eliminate plasmids, the cells were treated with mitomycin C, using the method of Rheinwald et al. [8].

For matings, the donor and recipient cells (at the approximate density of 10<sup>8</sup> cell ml<sup>-1</sup>) were mixed on LB plates in a ratio of 1:2. The plates were incubated for 2—4 h (in some cases up to 18 h) at 28 °C. The cells were scrapped off the plates and suspended in 0.56 % NaCl solution and transferred onto selective medium. The transconjugants obtained were passaged thrice on the respective media.

Bacterial strains used in this study

Strain (plasmid)	Relevant charac- teristics	Reference of Source
P. alcaligenes TR	w. t.,	Culture Collection, ICWC of the Ukrainian Acad. Sci.
P. testosteroni TO	w. t., Abs <sup>+</sup>	Ibid
P. fluorescence TR	w. t., Kat <sup>+</sup>	*
P. mendocina TO	w. t., Alk <sup>+</sup>	*
P. stutzeri AT	w. t., Etn <sup>+</sup>	*
P. putida TO	w. t., Abt <sup>+</sup>	*
P. desmolytica C37	w. t., Clm <sup>+</sup>	*
P. rathonis T	w. t., Vol <sup>+</sup>	*
Achromobacter eurydice TK	w. t., Dns <sup>+</sup>	*
P. putida K	w. t., Met <sup>+</sup>	*
Pseudomonas sp. 2T/1	w. t., Sds+	*
P. aeruginosa 1C	w. t., Las+	*

Note: w. t. — wild type or prototroph; utilization: Abs — alkyl benzenesulfonate; Kat — katamine; Alk — alkamone; Etn — etonium; Abt — amidobethaine; Clm — cyclimide; Abp — alkyl aminobispropionate; Vol — volgonat; Dns — disodium mono-2(alkanoamino)ethylsuccinatosulfonate; Las — alkylsulfate; Met — metaupon.

Several modifications of the alkaline lysis method were used to isolate plasmid DNA [9—13]. The size of plasmid DNA was estimated by comparing its electrophoretic mobility with standard plasmid DNAs. pLAFR5 (21.5 kb), RP4 (55 kb), NPL-1 (98 kb) and pBS2 (130 kb) which were used as marker plasmids.

The digestion of plasmid DNA was performed by *EcoRI* («Promega», USA) according to the supplier's protocol.

Results and Discussion. As the biodegradation of different synthetic compounds are frequently controlled on the genetic level by degradative plasmids (D-plasmids) [14, 15], the selected strains (Table) were tested for the presence of plasmid DNA. The most strains of bacteria-destructors have been selected under the continuous-flow conditions. Continuous cultivation was carried out in bioreactor of column type with a ceramic carrier under forced aeration. The immobilization of bacteria on a ceramic carrier was made by adhesion [4, 5].

A comparative study of the electrophoretic mobility of tested and marker plasmids of known sizes allowed us to establish the following: strains-degraders of anionic surfactants: P. rathonis T (Vol<sup>†</sup>), P. aeruginosa 1C (Las<sup>†</sup>), Pseudomonas sp. 2T/1 (Sds<sup>†</sup>), P. putida K (Met<sup>†</sup>) have one plasmid of about 100 kb; strain P. alcaligenes TR (ABS<sup>†</sup>) have one plasmid of about 130 kb; strain Achromobacter eurydice TK (Dns<sup>†</sup>) have one plasmid with less size (Fig. 1); strains P. fluorescens TR (Alk<sup>†</sup>), P. mendocina TO

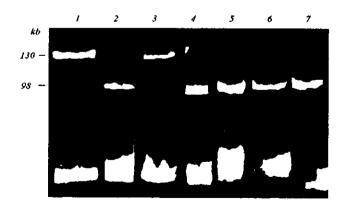


Fig. 1. Electrophoretic mobility of plasmids DNA of bacteria-degraders of anionic surfactants: I - Pseudomonas putida 231 (pBS2) (130 kb); 2 - P. putida 202 (NPL-I) (98 kb); 3 - P. alcaligenes TR; 4 - P. rathonis T; 5 - P. putida K; 6 - Achromobacter eurydice TK; 7 - P. aeruginosa 1C

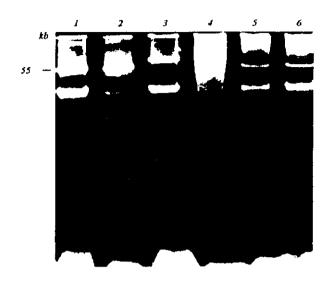


Fig. 2. Electrophoretic mobility of plasmids DNA from rains of bacteria-degraders of cationic and ampholytic surfactants: I—plasmid RP4 (55 kb); 2-P. fluorescence TR; 3-P. mendocina TO; 4, 5-P. putida TO; 6-P. stutzeri AT

(Etn<sup>+</sup>) harboured one plasmid each of about 60—70 kb. The electrophoretic mobility of the plasmids detected in strains *P. putida* TO (Abp<sup>+</sup>), *P. stutzeri* AT (Abt<sup>+</sup>) was close to the marker plasmid RP4 of 55 kb (Fig. 2).

We have failed to detect plasmid DNA in straindestructor of cyclimide, P. desmolytica C37, P. testosteroni TO, by screening techniques for realization of plasmid DNA such as alkaline lysis method with several modifications [10—13].

Fig. 3 demonstrates the fragments of plasmid DNA isolated from degrader strains digested by *EcoRI*. It has been shown that the different plasmids of bacteria-degraders have different restriction patterns.

Curing of plasmids may be induced by chemical reagents, i. e. acridine orange NO, ethidium bromide, SDS, mitomycin C (MC). For curing plasmid DNA was used MC in concentration  $1,25-30.0~\mu g \cdot ml^{-1}$ . The sublethal MC concentrations were  $20~\mu g \cdot ml^{-1}$  for anionic degrader — strain P. alcaligenes TR and strains degraders of ampholytic and cationic surfactants  $30.0~and~20.0~\mu g \cdot ml^{-1}$ , respectively. The clones which lost the ability for utilization of surfactants (Sac<sup>+</sup> phenotype) were obtained for strains P. rathonis T, P. alcaligenes TR, P. putida TO, P. stutzeri AT, P. fluorescens TR, P. putida K and P.

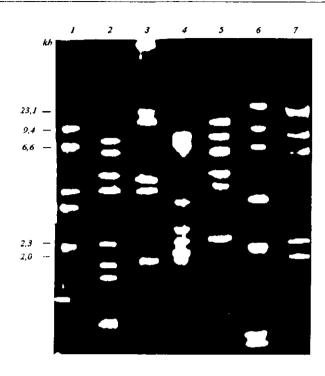


Fig. 3. Electrophoretic mobility of restrictions patterns DNA plasmids of strains-degraders of surfactants: I - P. mendocina TO; 2 - Achromobacter eurydice TK; 3 - P. putida TO; 4 - P. alcaligenes TR; 5 - P. fluorescence TR; 6 - P. rathonis T; 7 - P. marker DNA phage  $\lambda/H$ indIII

mendocina TO. The occurrence frequency of such variants was 0.4-1.8 %. We failed to observe the reversions to Sac<sup>+</sup> phenotype in most cured mutants. Plasmid DNA was not found in clones with lost ability for utilization of surfactants, except for the strain P. fluorescens TR.

The results obtained indicate the plasmid localization of the surfactant degradation determinants. To study whether the plasmids of strains degrading surfactants are able to conjugative transfer the experiments on mating have been carried out.

The transconjugants obtained were selected on the media supplemented with an appropriate surfactant as a sole carbon source. The plasmid DNA of expected molecular mass has appeared in transconjugant cells. The frequency of appearing Sac<sup>+</sup> phenotype was  $10^{-6}$  to  $10^{-3}$ .

The high frequency of the Alk<sup>+</sup> phenotype losing induced by mitomycin C favours the above suggestion. To elucidate the ability of the plasmid from strain P. fluorescens TR for conjugative and mobilization trans-

fer, we tested its ability to mobilize the non-conjugative plasmid pBS222 A-15. This plasmid was transferred to strain P. fluorescens TR by using a helper plasmid pRK2013 followed by a series of matings in which the above strain was employed as a donor of plasmid pBS222 A-15, while strains of E. coli and P. putida were employed as recipients. If own plasmid of strain P. fluorescens was conjugative, the mobilization transfer of a small plasmid pBS222 A-15 to recipient cells would be observed. Yet, numerous experiments failed to reveal this plasmid conjugative transfer at a frequency less than 10<sup>-9</sup>. So, the question related to the resident plasmid ability for conjugation still remains open. As for this plasmid functional significance, one may probably consider it to be the D-plasmid harbouring genes that control alkamone utilization.

Thus, we have shown that the degradation of anionic, cationic and ampholytic surfactants by pseudomonas is probably controlled by plasmid genes. Most plasmids found in strains-degraders of surfactants proved to be conjugative and may be used by bacteria for the degradation of ecologically hazardous compounds in the purification systems.

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Плазміди біодеградації поверхнево-активних речовин

### Резюме

Показано, що деградація багатьох аніонних, катіонних та амфолітних поверхнево-активних речовин (ПАР) у псевдомонад контролюються плазмідами розміром 60—130 тис. п. н. Більшість плазмід здатні до кон'югативного переносу та елімінації з бактеріальних клітин. Створені первинні рестрикційні карти плазмід біодеградації ПАР мають значну різницю.

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Плазмиды биодеградации поверхностно-активных веществ

## Резюме

Показано, что деградация многих анионных, катионных и амфолитных поверхностно-активных веществ у псевдомонад контролируется плазмидами размером 60—130 тыс. п. н. Большинство плазмид способны к конъюгативному переносу и элиминации из бактериальных клеток. Полученные первичные рестрикционные карты данных плазмид имеют значительные различия.

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