

Characterization of SAL plasmid isolated from *Pseudomonas putida*

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Pseudomonas putida 에서 분리한 SAL 플라스미드의 특성

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ABSTRACT: Three strains of bacteria utilizing salicylate, KU801(pKU5, pKU8), KU803 (pKU6, pKU9), and KU806(pKU7, pKU10), were selected from the isolates and identified as *Pseudomonas putida*. By agarose gel electrophoresis, it was found that the strains had two plasmids each. All three strains were resistant to antibiotics such as ampicillin, tetracyclin, and chloramphenicol, and did not utilize other aromatic and aliphatic hydrocarbons examined except salicylate. The plasmids(pKU5, pKU6, and pKU7) of larger molecular weight were cured by treatment with mitomycin C and frequencies of curing were 0.4%, 1.67%, and 0.75%, respectively. Cured strains did not degrade salicylate and still had antibiotic resistances, which were identical with wild strains. The genes for salicylate degradation were proved to be encoded on their plasmids. The molecular weights of pKU 5 and pKU 6 were estimated as 103.5 Md, and that of pKU 7 as 101 Md. The new SAL plasmids, pKU 5, pKU 6, and pKU 7 were transferred to *P. putida* and *P. aeruginosa*, but not to *E. coli*.
KEY WORDS □ SAL plasmid; *Pseudomonas putida*.

Many synthetic chemicals are added to our environment in the form of herbicides, pesticides and industrial effluents, and the problem of counteracting or neutralizing these wastes is becoming urgent. Soil microorganisms, particularly those belonging to the genus *Pseudomonas*, are able to transform or utilize a wide range of natural and synthetic organic compounds(Stanier *et al.*, 1966). Recent works have indicated that, in certain strain, the genes responsible for the degradative functions are carried not on the chromosome but are borne on plasmids. Thus the nutritional versatility can be achieved by the occurrence of

broadly transmissible plasmids(Benson and Shapiro, 1978). Conjugative extrachromosomal elements conferring metabolic specificity for the degradation of salicylic acid(SAL), naphthalene(NAH), and xylene and toluene(TOL) have been isolated and studied(Chakrabarty, 1972; Dunn and Gunsalus, 1973; Worsey and Williams, 1975). Nonconjugative degradative plasmids have been also described, including one specific for toluene and xylene(XYL; Friello *et al.*, 1976).

In this report, we describe on the properties of new SAL plasmids isolated from *Pseudomonas putida*.

MATERIALS AND METHODS

Bacterial strains and plasmids

Bacterial strains and plasmids used in this study are listed in Table 1.

Media and culture conditions

The medium L broth (Miller, 1972) was used as a complete medium, and M9 (Maniatis *et al.*, 1982) was used as a minimal medium. For growth of *Pseudomonas* strains utilizing salicylate as a sole source of carbon, the M9 medium containing 10 mM sodium salicylate instead of glucose (M9-Sal) was used. Incubations were carried out at 30°C for *P. putida* and at 37°C for *P. aeruginosa* and *E. coli*. For the carbon source utility test, cells were inoculated into M9 medium containing each aromatic hydrocarbon instead of glucose and incubated for one week at 30°C as previously (Kim and Lee, 1984). Used carbon sources

were salicylate, toluene, xylene, benzene, cyclohexane, naphthalene, camphor, octane, and catechol.

Isolation and identification of bacterial strains

Sampling was done from February to June in 1985. Soil samples of stream and industrial waste were taken from the areas thought to have been polluted in Seoul and its suburbs. The enrichment and isolation of bacterial strains utilizing salicylate as a sole source of carbon were carried out as described previously (Kim and Lee, 1984). The species of selected three strains having plasmids among isolates were identified. Identification test was carried out by the method of Palleroni (1984).

Isolation of plasmid DNA

Crude lysates of plasmid DNAs were prepared according to the method of Nakazawa *et al.* (1980). Cells were grown overnight in

Table 1. Bacterial strains and their characteristics.

Strain/Plasmid	Relevant characteristics	Source or Reference
<i>Pseudomonas putida</i>		
KU801 (pKU5, pKU8)	wild type/pKU5 (Sal ⁺)	This work
KU803 (pKU6, pKU9)	wild type/pKU6 (Sal ⁺)	This work
KU806 (pKU7, pKU10)	wild type/pKU7 (Sal ⁺)	This work
TN1032	<i>trp⁻ leu⁻ ben1⁻ Str^r</i>	Nakazawa and Yokota (1977)
TN1307	<i>trp⁻ leu⁻ ben1⁻ Str^r recA</i>	Nakazawa and Yokota (1977)
TN1126	<i>met⁻ trp⁻</i>	Nakazawa and Yokota (1977)
TN1126 (RSF1010)	<i>met⁻ trp⁻/RSF1010 (Sm^r)</i>	Transformation of TN1126 with RSF1010
<i>Pseudomonas aeruginosa</i>		
PA0303 (pMG1)	<i>arg⁻/pMG1 (Sm^r)</i>	Hansen and Olsen (1978)
PA0303 (Rms148)	<i>arg⁻/Rms148 (Sm^r)</i>	Sagai <i>et al.</i> (1975)
PA02003	<i>arg⁻ recA</i>	Olsen <i>et al.</i> (1982)
PA08 (R18)	<i>met⁻ ilv⁻ Str^r/R18 (Km^r)</i>	Issac and Holloway (1968)
KU815 (R18, pKU6, pKU9)	<i>/R18 (Km^r) pKU6 (Sal⁺)</i>	Conjugation of PA08 with KU803
<i>Pseudomonas</i> sp		
KU802, KU804, KU805 KU807, KU808	wild type (Sal ⁺)	This work
<i>Escherichia coli</i>		
HB101	<i>r⁻ m⁻ pro⁻ leu⁻ Str^r recA</i>	Hamer <i>et al.</i> (1976)
HB101 (pRK290)	<i>/pRK290 (Tc^r)</i>	Corbin <i>et al.</i> (1982)
C600 (RP4)	<i>/RP4 (Ap^r Tc^r Km^r)</i>	Barth and Grinter (1977)

L broth. Harvested cells were treated with lysozyme (1mg/ml) in 25% Sucrose-50 mM Tris-HCl(pH 8.0) for 5 min at 0°C, followed by incubation for another 5 min in the presence of 6.25 mM EDTA. The cells then treated with 6.25 mM EDTA. The cells then treated with 0.5% Brji 58-0.2% sodium deoxycholate-62.5 mM EDTA in 50 mM Tris-HCl(pH 8.0) for 10 min at 0°C, and centrifuged at 30,000×g for 20 min at 2°C. Plasmid DNA contained in the supernatant was precipitated with 10% polyethylene glycol in the presence of 0.5 M NaCl. The precipitate was collected by centrifugation and dissolved in 50 mM Tris-HCl-5 mM EDTA-5 mM NaCl(pH 8.0) and then subjected to electrophoresis.

Agarose gel electrophoresis

Agarose gel electrophoresis was performed in a vertical slab gel of 0.7% agarose in TAE buffer composed of 0.4 M Tris, 0.2 M sodium acetate and 0.01 M EDTA at pH 8.0. Gels were run at 120 V for 2 hr. Gels were stained in a solution of ethidium bromide (1 µg/ml) for 20 min, rinsed and photographed under UV illumination.

Antibiotics resistance test

For the determination of resistance to various antibiotics, bacterial suspensions were plated on L agar containing different concentrations of various antibiotics respectively and incubated for 48 hr at 30°C.

Curing with mitomycin C

The method of Rheinwald *et al.* (1973) was used as follows. Cells were inoculated into 5 ml of L broth. After overnight growth 0.1 ml of the culture (10^4 to 10^5) were inoculated into 5 ml of L broth containing mitomycin C and incubated with shaking for 48 hr. The culture was then diluted and plated on L agar. Colonies were then replica-plated to another L agar plate and also M9-Sal agar plate.

Bacterial conjugation

Transfer frequency was determined by mating on membrane filters as described by De Graaf *et al.* (1973). Exponentially growing donors were mixed with an equal volume of

recipient culture in the exponential phase. The mixture was filtered on a membrane filter (0.2 µm pore size, 25 mm diameter Gelman Science, INC). Membrane was placed on the surface of a freshly prepared Penassay broth agar plate and incubated for 5 hr at 30°C or 37°C. Bacteria were then suspended in 1 ml of saline, and 0.1 ml samples of suitable dilutions were spread on the selective medium. Conjugation frequency was determined after the titration of donor cells on L agar plates. When *P. aeruginosa* PAO strains were used as recipients, recipient cells were heated prior to mating for 10 min at 50°C (Curtiss III, 1981).

RESULTS AND DISCUSSION

Isolation and identification of bacterial strains

Eighty-six bacteria strains utilizing salicylate as a sole source of carbon were isolated. Three strains, KU801, KU803, and KU806, which contained plasmids respectively, were selected among them (Fig.1). From the morphological, physiological, and biochemical characteristics of the strains, they were identified as *Pseudomonas putida* respectively.

Carbon source utility

Selected strains, KU801, KU803, and KU806, were utilized salicylate and catechol as a sole source of carbon. Toluene, xylene, benzene, cyclohexane, naphthalene, camphor, and octane were not utilized by the selected strains.

Resistance to antibiotics

The resistances of the selected strains to different concentration of various antibiotics tested were very similar. All the three strains had resistance to 1600 µg/ml of chloramphenicol, 800 µg/ml of ampicillin, 50 µg/l of tetracyclin, but were sensitive to more than 25 µg/ml of gentamycin, streptomycin, and kanamycin respectively.

Curing of cells harbouring SAL plasmid

When KU801, KU803, and KU806 were

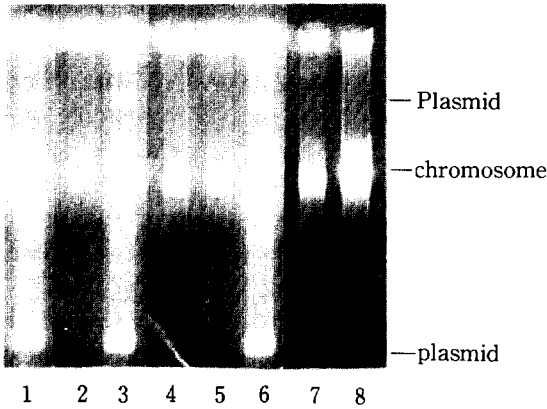


Fig.1. Agarose gel electrophoresis of plasmids isolated from *Pseudomonas* strains.

1. *P. putida* KU801
2. *Pseudomonas* sp. KU802
3. *P. putida* KU803
4. *Pseudomonas* sp. KU804
5. *Pseudomonas* sp. KU805
6. *P. putida* KU806
7. *Pseudomonas* sp. KU807
8. *Pseudomonas* sp. KU808

* The DNA bands between chromosome and small plasmid in lane 1, 3, and 6 are oc form of small plasmid.

treated with various concentration of mitomycin C, no growth appeared with 17.5 $\mu\text{g/ml}$, 12.5 $\mu\text{g/ml}$, and 20 $\mu\text{g/ml}$, respectively and thus 15 $\mu\text{g/ml}$, 10 $\mu\text{g/ml}$, and 17.5 $\mu\text{g/ml}$ were considered as sublethal concentration in KU801, KU803, and KU806, respectively. Loss of the ability to grow on salicylate at these mitomycin C concentration was varied from 0.40% to 1.67% of total viable colonies (Table 2). Agarose gel electrophoresis of the cured strains (Fig.2) also indicates that the inability to utilize salicylate is due to the loss of

Table 2. Curing frequency of the selected *Pseudomonas putida* strains by mitomycin C.

Strains	Concentration of mitomycin C ($\mu\text{g/ml}$)	No. of colonies tested	Used marker for curing	Frequency of curing (%)
KU801	15	500	Salicylate	0.40
KU803	10	900	Salicylate	1.67
KU806	17.5	936	Salicylate	0.75

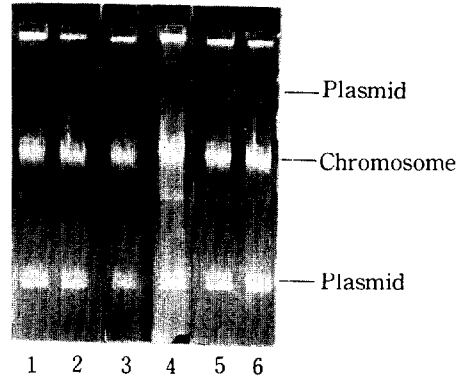


Fig.2. Agarose gel electrophoresis of the cured strains.

1. *P. putida* KU801
2. *P. putida* KU811 (Cured strain of KU801)
3. *P. putida* KU803
4. *P. putida* KU813 (Cured strain of KU803)
5. *P. putida* KU806
6. *P. putida* KU816 (Cured strain of KU806)

plasmid-borne genes which involves in salicylate degradation. Plasmids of the small molecular weight still exist in the cured strains and antibiotic resistances of cured strains are identical with those of the native strains. It was suggested that the small plasmids were nothing to do with salicylate degradation, but with antibiotic resistance.

Conjugative transfer of the ability of salicylate degradation

The fact that the ability of salicylate degradation is plasmid-borne is also inferred from the data in Table 3. SAL plasmids can be transferred at a frequency of 10^{-5} to 10^{-7} into *P. putida* and 10^{-6} to 10^{-7} into *P. aeruginosa*. It has been reported that SAL plasmid undergoes restriction when *P. aeruginosa* PAO strains is grown at 30°C or 37°C (Chakrabarty, 1972; Kanemitsu, 1980). However, when *P. aeruginosa* PAO strains were heated and then used as recipients (in Methods), transfer of SAL plasmid occurred without restriction. The SAL plasmid in *P. aeruginosa* KU815, conjugant formed by mating PAO8 as a recipient with KU803 as a

Table 3. Conjugative transfer of the ability of salicylate degradation to various strains.

Donor	Recipient	Selected marker**	Frequency of transfer*
<i>P. putida</i> KU801	<i>P. putida</i> TN1032	<i>trp, leu, Sm, Sal</i>	4.0×10^{-5}
	<i>P. putida</i> TN1307	<i>trp, leu, Sm, Sal</i>	3.0×10^{-6}
	<i>P. aeruginosa</i> PAO2003	<i>arg, Sm, Sal</i>	1.8×10^{-6}
	<i>P. aeruginosa</i> PAO8	<i>met, ilv, Sm, Sal</i>	1.0×10^{-6}
	<i>E. coli</i> HB101	<i>pro, leu, Sm, Sal</i>	0
<i>P. putida</i> KU803	<i>P. putida</i> TN1032	<i>trp, leu, Sm, Sal</i>	2.0×10^{-6}
	<i>P. putida</i> TN1307	<i>trp, leu, Sm, Sal</i>	3.0×10^{-7}
	<i>P. aeruginosa</i> PAO2003	<i>arg, Sm, Sal</i>	5.0×10^{-7}
	<i>P. aeruginosa</i> PAO8	<i>met, ilv, Sm, Sal</i>	1.6×10^{-6}
	<i>E. coli</i> HB101	<i>pro, leu, Sm, Sal</i>	0
<i>P. putida</i> KU806	<i>P. putida</i> TN1032	<i>trp, leu, Sm, Sal</i>	2.0×10^{-5}
	<i>P. putida</i> TN1307	<i>trp, leu, Sm, Sal</i>	1.0×10^{-6}
	<i>P. aeruginosa</i> PAO2003	<i>arg, Sm, Sal</i>	9.0×10^{-8}
	<i>P. aeruginosa</i> PAO8	<i>met, ilv, Sm, Sal</i>	2.0×10^{-6}
	<i>E. coli</i> HB101	<i>pro, leu, Sm, Sal</i>	0
<i>P. aeruginosa</i> KU815	<i>P. putida</i> TN1307	<i>trp, leu, Sal</i>	4.1×10^{-7}

* Frequency of transfer: number of conjugants/number of donors

** The media used were M9 minimal media.

donor, could be transferred at a frequency of 4.1×10^{-7} back into *P. putida*. These results indicated that SAL plasmids which isolated in this study are not restricted in *P. aeruginosa* PAO strain.

Agarose gel electrophoresis of the conjugant strains (Fig.3) shows that both of the plasmids in donor cells were transferred to the recipient cells. This seems to be a result that transmissible SAL plasmid provides the transfer to nontransmissible small plasmid (Freifelder, 1983). When SAL plasmid was transferred to *E. coli*, conjugants were not obtained. However, it is not clear whether the transmissible plasmids is not transferred to the members of other genera (Chakrabarty, 1972, Chakrabarty, 1976) or in enteric bacteria were not expressed the genes for hydrocarbon oxidation (Farrell and Chakrabarty, 1979).

Determination of molecular weight of plasmid DNA

The molecular weight of the plasmids was determined using covalently closed circular

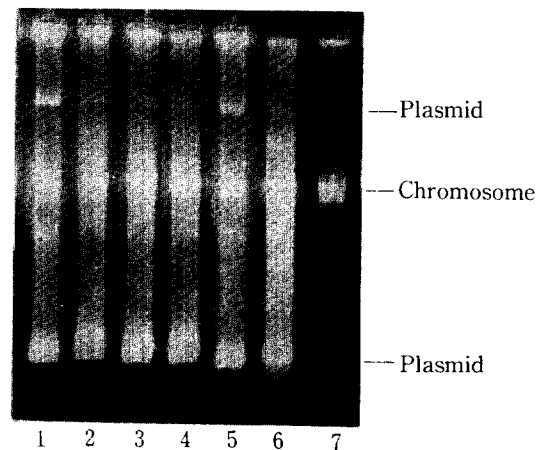


Fig.3. Agarose gel electrophoresis of the conjugants.

1. *P. putida* KU801 (donor)
2. *P. putida* KU812 (conjugant of TN1307 with KU801)
3. *P. putida* KU803 (donor)
4. *P. putida* KU814 (conjugant of TN1307 with KU803)
5. *P. putida* KU806 (donor)
6. *P. putida* KU817 (conjugant of TN1307 with KU806)
7. *P. putida* TN1307 (recipient)

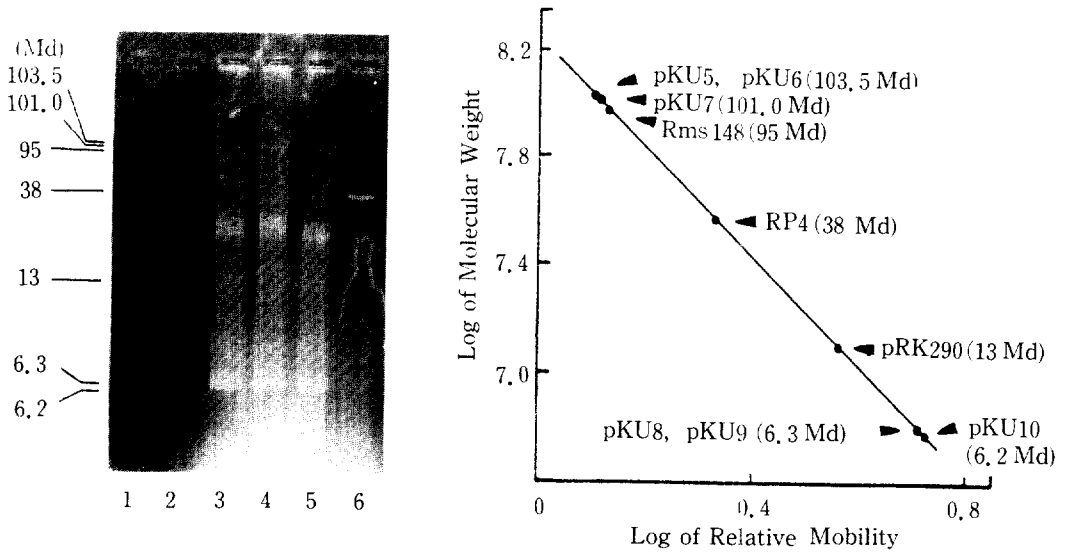


Fig.4. Molecular weight of plasmids.

1. pRK290 2. Rms148 3. pKU5, pKU8 4. pKU6, pKU9 5. pKU7, pKU10 6. RP4

forms of pRK290 (13 Md), RP4 (38 Md), Rms 148 (95 Md) as standard size markers (Fig.4). The SAL plasmids of KU801, KU803, and KU806 were named pKU5, pKU6, and pKU7, and their small plasmids, pKU8, pKU9, and pKU10, respectively. The molecular weights of pKU5 and pKU6 were estimated as 103.5 Md, and that of pKU7 as 101

Md. Their sizes are rather large in comparison with 40 Md (Palchudhuri and Chakrabarty, 1976), 43 Md (Johnston *et al.*, 1977), 45 Md (Gunsalus *et al.*, 1981), 48 Md (Heinaru *et al.*, 1978), and 55 Md (Farrell *et al.*, 1978) which have been previously reported. Therefore, SAL plasmids examined in this research seem to be new SAL plasmids.

적 요

분리한 살리실산 자화세균 중 플라스미드를 갖는 세 균주를 선별하였다. 세 균주, KU801(pKU5, pKU8), KU803(pKU6, pKU9), KU806(pKU7, pKU10)는 각각 두 개씩의 플라스미드를 가지고 있음이 전기영동에 의해 밝혀졌고, *Pseudomonas putida*로 동정되었다. 세 균주들은 모두 암피실린, 테트라사이클린, 클로람페니콜등의 항생제에 대하여 내성을 지니며, 조사된 방향족과 지방족 탄화수소들 중 살리실산과 그의 중간 대사물인 카테콜만을 이용하였다. 큰 분자량의 플라스미드 (pKU5, pKU6, pKU7)는 마이토마이신 C로 처리하였을 때 큐어되며 그 빈도는 각각 0.40%, 1.67%, 0.75% 이었다. 큐어된 균주는 살리실산을 분해하지 못하였으나, 여전히 야생균주와 동일한 항생제 내성을 가지고 있었다. 살리실산 분해에 관여하는 유전자가 그들 플라스미드에 있는 것으로 판명되었다. pKU5와 pKU6의 분자량은 103.5Md, pKU7의 분자량은 101 Md으로 측정되었다. SAL 플라스미드인 pKU5, pKU6, pKU7은 접합에 의해 *P. putida*와 *P. aeruginosa*로는 전달되었으나, *E. coli*에서는 발현되지 아니하였다.

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