

CUPRIC ION RESISTANCE AS A NEW GENETIC MARKER OF A TEMPERATURE

SENSITIVE R PLASMID, Rts1 IN ESCHERICHIA COLI

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SUMMARY. A temperature sensitive kanamycin (Km) resistant R plasmid, Rts1, was found to confer cupric ion (Cu^{2+}) resistance on its hosts in Escherichia coli. At conjugal transfer, two kinds of segregants were obtained from Rts1, i.e. Cu^{2+} resistant, Km sensitive and Km resistant, Cu^{2+} sensitive plasmids. Protein T existed in E. coli cells harboring Rts1 or the $\text{Cu}^{\text{r}}\text{Km}^{\text{s}}$ -plasmid. The inhibitory effect on the host cell growth at 43°C was observed with Rts1⁺ or the $\text{Km}^{\text{r}}\text{Cu}^{\text{s}}$ -plasmid⁺ cells. A relationship between these Rts1 derivatives and Rts1 in Proteus mirabilis which has been studied was discussed.

Rts1 is a kanamycin (Km) resistant R plasmid and its replication and conjugal transfer are temperature sensitive (1,2). In addition, Rts1 inhibits the host cell growth at 42°C (3,4). Our recent studies revealed the presence of protein T, a temperature sensitive protein, in the outer membrane of the Rts1⁺ cells (5), and the production of extracellular DNase from the Rts1⁺ cells (6) of Escherichia coli strains. We are intending to find the additional genetic markers other than Km on Rts1 genome which would allow to select easily the Rts1⁺ cells. As a result we found a resistance to cupric ion (Cu^{2+}) on Rts1.

MATERIALS AND METHODS

Bacterial strains and R plasmids. Bacterial strains and R plasmids used were shown in Table 1.

Media for isolation and selection of Rts1⁺ cells. M-9 glucose agar supplemented with amino acids (20 $\mu\text{g}/\text{ml}$) and thiamine (2 $\mu\text{g}/\text{ml}$) if required, was used as basal medium. When Km or Cu^{2+} resistance was transferred from XA7012 (Rts1) to W677, 0.1ml of the mixed culture was plated on the M-9 agar containing nalidixic acid (Nal: 50 $\mu\text{g}/\text{ml}$) and either Km (50 $\mu\text{g}/\text{ml}$) or CuSO_4 (1mM). In the mating between W677 R⁺ and JC1569, streptomycin (100 $\mu\text{g}/\text{ml}$) was substituted for Nal.

Analysis of plasmid DNA in alkaline sucrose gradient. Cell lysate was prepared from the culture of JC1569 R⁺ cells labeled with 40 $\mu\text{Ci}/\text{ml}$ of ^3H -thymidine by the procedure of Freifelder et al. (7). The lysate was layered onto a 5ml 5 to 20% linear alkaline sucrose gradient containing 0.5M NaCl, 0.02M EDTA and 0.3M NaOH, and centrifuged at 40,000rpm in a Hitachi RPS65T rotor for 18 min at 20°C . The fractions were collected from the bottom onto filter paper, washed with 5% trichloroacetic acid and counted in a Packard liquid scintillation counter.

Isolation of outer membrane. Outer membrane was prepared by the methods as described previously (5).

Polyacrylamide gel electrophoresis. SDS-PAGE and two-dimensional gel electrophoresis were carried out as described previously (5) except the condition of the solubilization of the samples.

RESULTS

Transfer of Km and Cu²⁺ resistances. Stock culture of XA7012(Rts1) in Pennasay broth (Difco, PAB) was found to be composed of Km^rCu^r, Cu^r and Km^r cells, by the ratio of 75, 10 and 15%, respectively. The minimal inhibitory concentrations of CuSO₄ to XA7012 R⁻, Km^r and Cu^r cells were 0.06mM, 0.06mM and 10mM, on the M-9 glucose plates. This finding led us to examine the transferability of Cu²⁺ resistance. XA7012(Rts1) resistant both to Km and Cu²⁺ was mated with W677 Nal^r at 30°C for 17h. The transconjugants were selected on the plates containing Nal and either Km or CuSO₄. W677 cells that received Cu²⁺ resistance formed yellowish brown colonies on the plate containing CuSO₄, whereas the Cu²⁺ sensitive cells could not grow on the plate. When Km was used for the selection, only 2% of transconjugants were resistant both to Km and Cu²⁺. In contrast, when the selection was made for the Cu²⁺ resistance, 97% of transconjugants received the both resistances. Thus we obtained three types of Rts1 plasmids at conjugal transfer, i.e. Rts1(KmCu), Rts1(Cu) and Rts1(Km). These three plasmids were transferred further from W677 to JC1569, and their integrity as plasmid was examined in this recA⁻ host as follows.

Covalently closed circular (CCC) plasmid DNA. DNAs of JC1569 strains having either one of the plasmids were analyzed by alkaline sucrose density gradient centrifugation (Fig.1). Each cell lysate showed single peak which should be CCC-DNA of the respective plasmid. By cosedimentation studies with R28K (44x10⁶ daltons) or NR1 (63x10⁶ daltons), the molecular sizes of Rts1(KmCu), Rts1(Cu) and Rts1(Km) were estimated to be approximately 140x10⁶, 120x10⁶ and 100x10⁶ daltons., respectively.

Host cell growth at the non-permissive temperature. JC1569 cells with or without the plasmid were grown to mid-exponential phase at 30°C. Then the cells were diluted 100 fold into the fresh medium and continued to grow at ei-

Table 1. Bacterial strains and R plasmids

Strain	Genetic character	source
Bacterium		
<i>E. coli</i> K-12		
XA7012	F ⁻ <u>Δlac galE</u>	J. R. Beckwith
W677	F ⁻ <u>lac leu thr thi</u> Nal ^r	laboratory strain
JC1569	F ⁻ <u>recA1 gal leu his</u> Str ^r	H. Uchida
W3104	F ⁻ <u>galT12</u>	laboratory strain
R plasmid		
Rts1(KmCu)	Km ^r Cu ²⁺ r	this study
Rts1(Cu)	Km ^s Cu ²⁺ r	this study
Rts1(Km)	Km ^r Cu ²⁺ s	this study

Km: kanamycin, Cu²⁺: cupric ion, Nal: nalidixic acid, Str: streptomycin
r: resistant, s: sensitive

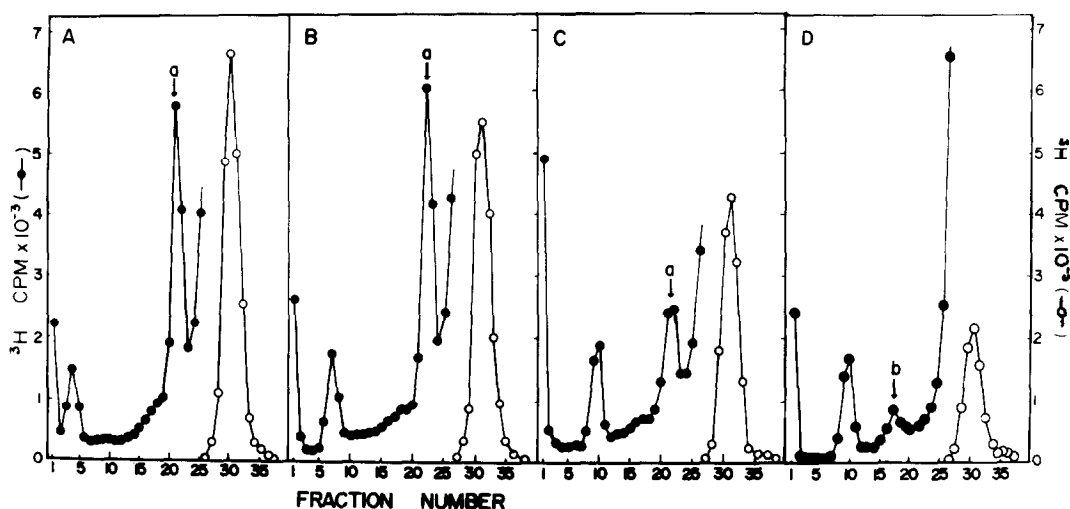


Fig.1 Alkaline sucrose gradient centrifugation of DNA from *E. coli* JC1569 (Rts1). Cells were grown in PAB containing 40uCi/ml of ³H-thymidine and 250ug/ml of deoxyadenosine for at least two generations at 30°C. Cell lysates were prepared and analyzed in 5-20% alkaline sucrose gradients, as described in MATERIALS AND METHODS. In each centrifugation, lysate from *E. coli* W3104 (R28K) or W3104(NR1) was cosedimented with Rts1⁺ cell lysate as a reference. (A) Lysates from JC1569(Rts1(KmCu)) and W3104(R28K). (B) Lysates from JC1569 (Rts1(Cu)) and W3104(R28K). (C) Lysates from JC1569(Rts1(Km)) and W3104(R28K). (D) Lysates from JC1569(Rts1(Km)) and W3104(NR1). In the figure, a and b indicate CCC-DNA of R28K and NR1, respectively. Note the changes of scale at fractions 26-27. The large peak on the right (o-o) is sheared DNA of the host chromosome.

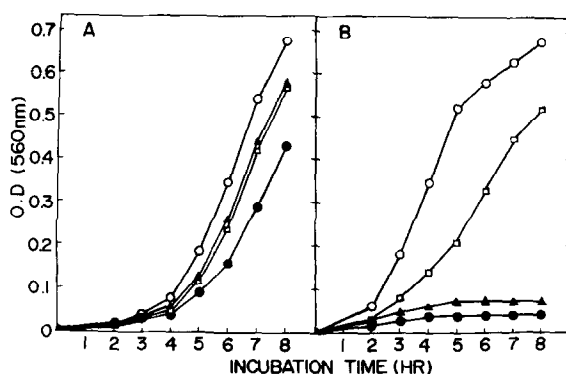


Fig.2. Growth of JC1569 R⁻, Rts1(Km)⁺, Rts1(Cu)⁺ and Rts1(KmCu)⁺ cells. The cells were grown to mid-exponential phase with shaking in PAB at 30°C. Each culture was diluted 100 fold by fresh PAB and continued to grow at 30°C (A) and 43°C (B). Bacterial growth was followed by measuring optical density of the culture at 560nm with Coleman Junior II spectrophotometer. Symbols:▲,JC1569 (Rts1(Km)),□,JC1569(Rts1(Cu)),●,JC1569(Rts1(KmCu)) and ○,JC1569 R⁻.

ther 30°C or 43°C. As shown in Fig.2, although the growth of Rts1(KmCu)⁺ and Rts1(Km)⁺ cells were remarkably inhibited at 43°C, Rts1(Cu) did not inhibit its host cell growth. Throughout these experiments the plasmid-cured cells were not detected at either 30°C or 43°C. Similar results were obtained in W677.

Protein T and other outer membrane proteins. Outer membrane proteins of XA7012, W677 and JC1569 cells with or without the plasmid were analyzed by SDS-PAGE (Fig.3). The W677 cells were analyzed further by two-dimensional gel electrophoresis (Fig.4). Protein T was found in all samples of the cells having Rts1(KmCu) or Rts1(Cu) (Fig.3). In regard to the "major proteins" (8) there was no difference among the four strains in W677 (Fig.4).

DISCUSSION

To our knowledge, this is the first report on plasmid mediated Cu²⁺ resistance in bacteria. Mechanisms of Cu²⁺ resistance has been reported previously in *E. coli* and yeast. In the former the resistance is ascribed to the disappearance of major protein b and c from the outer membrane, which inhibits penetration of the low molecular hydrophilic substances, including Cu²⁺ (9). In the latter, the Cu²⁺ resistance is due to production of a large amount of H₂S

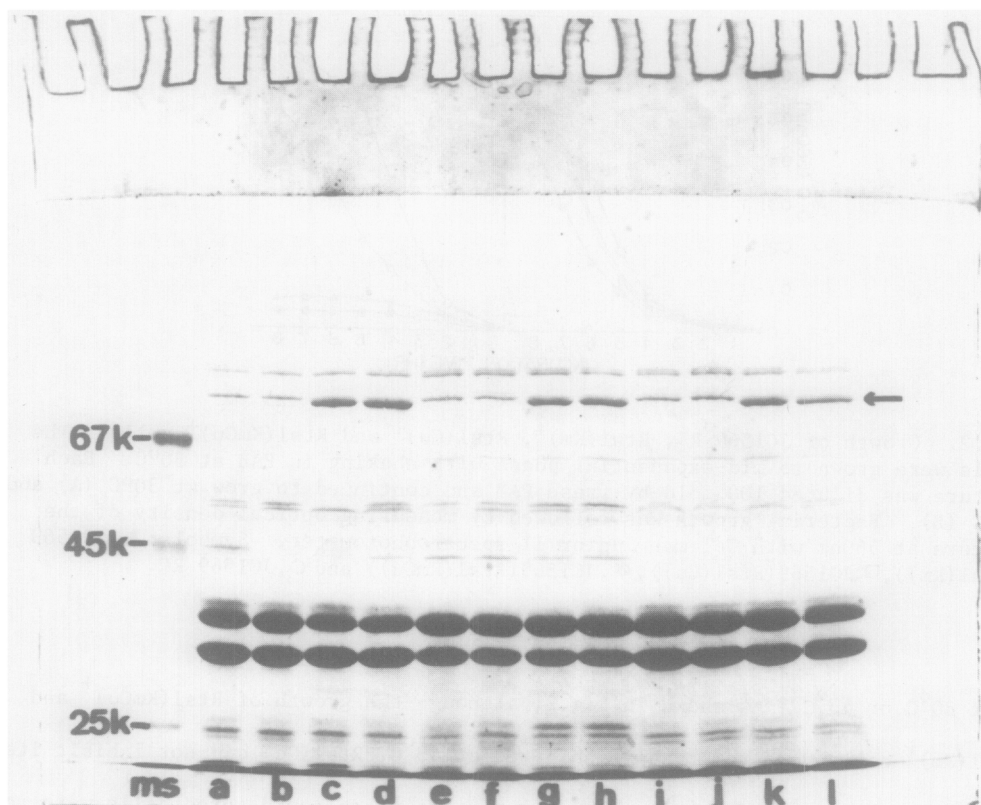


Fig.3. SDS polyacrylamide slab gel of outer membrane proteins from XA7012, W677 and JC1569 having either Rts1(Km), Rts1(Cu) or Rts1(KmCu), or without R plasmid. Purified outer membrane was solubilized by heating at 100°C for 3.5 min in "sampling buffer" (5), and applied to acrylamide slab gel (9%), containing 0.1% SDS. Molecular weight standards were: Bovine serum albumin (MW 67,000), hen egg albumin (MW 45,000) and chymotrypsinogen (MW 25,000). (a) XA7012 R⁻, (b) XA7012(Rts1(Km)), (c) XA7012(Rts1(Cu)), (d) XA7012(Rts1(KmCu)), (e) W677 R⁻, (f) W677(Rts1(Km)), (g) W677(Rts1(Cu)), (h) W677(Rts1(KmCu)), (i) JC1569 R⁻, (j) JC1569(Rts1(Km)), (k) JC1569(Rts1(Cu)), (l) JC1569(Rts1(KmCu)), (ms) molecular weight standards. —→, Protein T.

that results in formation of harmless CuS. In addition, the resistance in yeast is lost when grown without Cu²⁺ (10). However the Cu²⁺ resistance by Rts1(KmCu) and Rts1(Cu) was stably maintained regardless of presence or absence of Cu²⁺, and these cells resistant to Cu²⁺ had complete four major proteins in the outer membrane (Fig.4). Moreover, H₂S was not detected in the Rts1⁺ cells when tested in sulfide-indole-motility medium (data not shown). These findings suggest that the Cu²⁺ resistance of Rts1 might be due to a different mechanism from those described above.

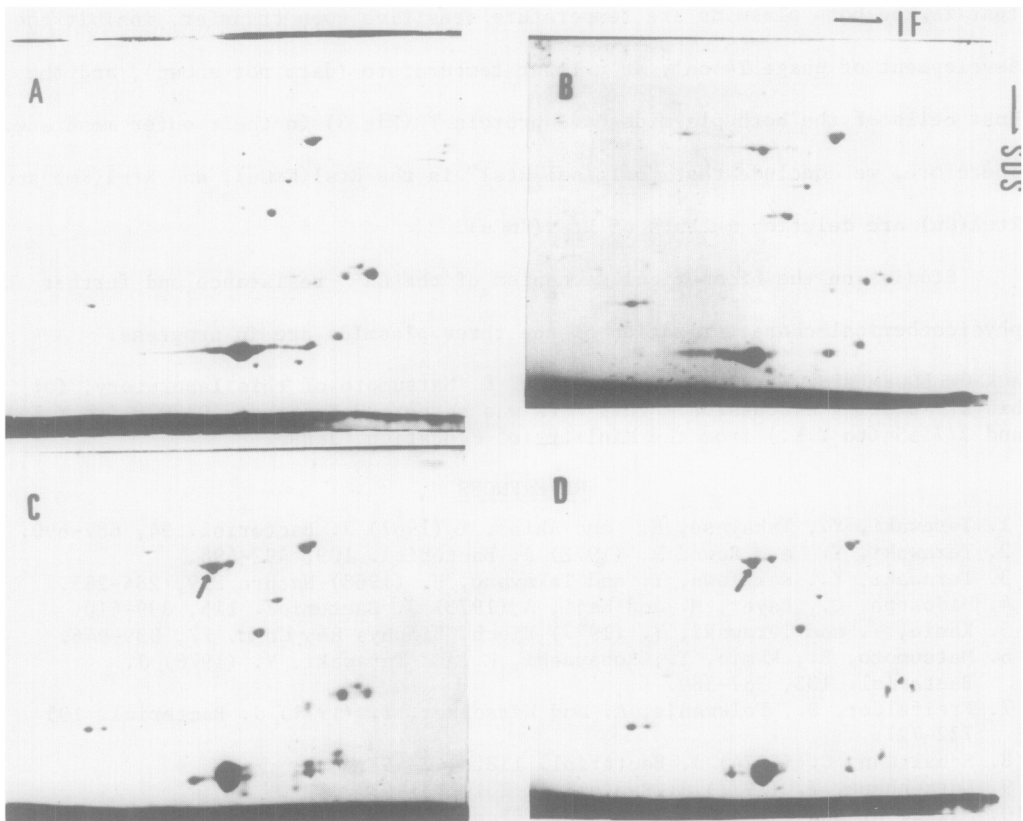


Fig.4. Two-dimensional slab gels of outer membrane proteins from W677 R⁻, Rts1(Km)⁺, Rts1(Cu)⁺ and Rts1(KmCu)⁺ cells.
 (A) W677 R⁻, (B) W677(Rts1(Km)), (C) W677(Rts1(Cu)), (D) W677(Rts1(KmCu)).
 —→, Protein T.

As shown in Fig.1, the presence of three kinds of Rts1 molecules were confirmed in *E. coli*. This finding raises following two questions; i) Which molecule is the "usual Rts1" that has been studied in this laboratory for long? ii) Was Rts1(Cu) molecule really derived from "original Rts1"? We have extensively studied the replication of Rts1 in *Proteus mirabilis* Pm17 (2,11,12). By a cosedimentation analysis of Rts1 in Pm17 and Rts1(Km) in JC1569 in alkaline sucrose gradient centrifugation, the both plasmids were found to sediment at the same position, indicating that "usual Rts1" corresponds to Rts1(Km) plasmid. As to the second question, Rts1(Cu) is considered to be a derivative of "original Rts1", since Rts1(Cu) shares several unusual properties in common with Rts1(KmCu),

that is, the both plasmids are temperature sensitive upon transfer, inhibit the development of phage T4 only at a lower temperature (data not shown), and the host cells of the both plasmids have protein T (Fig.3) in their outer membrane. Therefore, we conclude that "original Rts1" is the Rts1(KmCu), and Rts1(Km) and Rts1(Cu) are deletion mutants of Rts1(KmCu).

Studies on the biochemical mechanism of the Cu^{2+} resistance and further physicochemical characterization of the three plasmids are in progress.

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