# Isolation of Plasmid Deoxyribonucleic Acid from Two Strains of Bacteroides

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Two clinical isolates of Bacteroides contained covalently closed circular deoxyribonucleic acid (DNA) as shown by sedimentation in an alkaline sucrose gradient, CsCl ethidium bromide equilibrium centrifugation, and electron microscopy. Bacteriodes fragilis N1175 contained a homogeneous species of plasmid DNA with a molecular weight of  $25 \times 10^8$ . Bacteroides ochraceus 2228 contained two distinct, covalently closed circular DNA elements. The larger cosedimented with the covalently closed circular DNA form of the R plasmid, R100, corresponding to a molecular weight of  $70 \times 10^{\circ}$ ; the smaller sedimented as a 58S molecule with a calculated molecular weight of  $25 \times 10^6$ . The roles of these plasmids are unknown. Neither strain transferred antibiotic resistance to plasmid-negative Bacteroides or Escherichia coli, and neither produced bacteriocins active against other Bacteroides or sensitive indicator strains of E. coli.

Plasmids are stable, extrachromosomal, genetic elements which control many important bacterial properties (26) including conjugation, resistance to antimicrobials and heavy metals, bacteriocin production, and certain determinants of pathogenicity (23, 27). Because of the importance of several of these characteristics to human disease and microbial ecology, plasmids have been sought and found in a wide variety of bacteria including Pseudomonas (20), Bacillus (4, 24), Staphylococcus (26), Streptococcus (6, 7, 9, 15), and most genera of Enterobacteriaceae (7, 18, 26). These extrachromosomal elements usually exist in the bacterial cell as covalently closed circular deoxyribonucleic acid (DNA) molecules, but the conditions of extraction may result in considerable conversion to the open circular form (7, 18, 26).

Because Bacteroides spp. are among the most abundant members of the normal flora of human beings and would be logical reservoirs for R factors and bacteriocins, we were surprised that attempts to transfer antibiotic resistance from Bacteroides to sensitive Enterobacteriaceae or other Bacteroides were unsuccessful (1). When our attempts to transfer antibiotic resistance also failed, we extracted and studied the DNA from <sup>10</sup> strains of Bacteroides isolated from human infections. In this report we describe the isolation and characterization of

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covalently closed circular plasmid DNA from two of eight strains of tetracycline-resistant Bacteroides.

## MATERIALS AND METHODS

Bacteria. All strains of Bacteroides were isolated from human infections by the microbiology laboratory at University Hospital, University of California at San Diego, using an anaerobic system that has been described previously (11). The isolates were identified by standard biochemical methods and gas-liquid chromatography (19). Minimal inhibitory concentrations of penicillin, tetracycline, chloramphenicol, erythromycin, clindamycin, and rifampin were determined on agar plates of 5% rabbit blood in blood agar base (Difco Laboratories) containing serial twofold dilutions of antibiotic from 20 to 0.3  $\mu$ g/ml. Plates were incubated in GasPak jars (BBL). Strains of Bacteroides were screened for penicillinase production by a standard technique (28).

The restriction-negative Escherichia coli C and the plasmids R-100 and ColEl were generously provided by Don Helinski, Department of Biology, University of California at San Diego. E. coli 53E and Enterobacter cloacae 122H are R factor-negative recipients orginally isolated from an antibiotic-free population in Borneo (10). Enterobacter aerogenes K3151, a known producer of colicin (3), was used as a positive control in experiments testing for bacteriocin production. E. coli C and E. coli K-12 were used as colicin-sensitive indicators.

Radioactive labeling. The cells were diluted from an early stationary-phase culture into fresh Trypticase soy broth (Difco). After incubation for 6 h at 37 C, [methyl-3H]thymidine (0.01 mCi/ml; specific activity, 55.6 Ci/mmol; New England Nuclear Corp.) was added, and the cells were grown at 37 C to early stationary phase.

Cell lysis and sucrose gradient and CsCl-ethidium bromide centrifugation. Lysozyme spheroplasts were prepared, lysed with sodium dodecyl sulfate, and treated with NaCl to precipitate the chromosome as described by Guerry et al. (17). For sucrose gradient centrifugation, the lysate was diluted with 0.05 M tris(hydroxymethyl)aminomethane, pH 8.0, 0.05 M NaCl, and 0.005 M ethylenediaminetetraacetic acid, layered onto a 5 to 20% linear sucrose gradient in 0.05 M tris(hydroxymethyl)aminomethane, pH 8.0, 0.5 M NaCl, and 0.005 M ethylenediaminetetraacetic acid, and centrifuged in the Spinco SW50.1 rotor at 50,000 rpm for 55 min at 15 C. CsCl-ethidium bromide centrifugation, collection, scintillation counting, pooling, and dialysis of samples were performed as described by Helinski and co-workers (2, 5). Greater than 95% recovery of input counts was obtained from all velocity sedimentation runs.

Alkaline sucrose sedimentation. Samples pooled from the neutral 5 to 20% sucrose gradient were layered directly onto 20 to 31% alkaline sucrose gradients containing 0.3 N NaOH, 1.0 M NaCl, and 0.005 M ethylenediaminetetraacetic acid. Centrifugation in the Spinco SW50.1 rotor was for 55 min at 50,000 rpm and 15 C.

Electron microscopy. DNA samples, pooled from a preparative 5 to 20% neutral sucrose gradient, were prepared for electron microscopy by a modification of the Kleinschmidt procedure, as described by Davis et al. (12).

Conjugation experiments. These experiments were designed to detect transfer of antibiotic resistance from either tetracycline-resistant or multiply resistant Bacteroides fragilis to E. coli C, E. coli 53E, Enterobactor cloacae 122H, Bacteroides melaninogenicus, or to tetracycline-sensitive strains of B. fragilis. The general techniques for detection of transferable drug resistance (10, 31) were modified as appropriate for an anaerobic system. Stationaryphase cultures were prepared in brain heart infusion broth incubated in a GasPak jar, and 0.5 ml of the resistant strain was mixed with 3.5 ml of the sensitive strain. After overnight incubation, 0.5 ml of the mixture was inoculated over the surface of blood agar plates with and without antibiotics. When B. fragilis was the prospective donor and E. coli or Enterobacter cloacae the recipient, the inoculum was taken from the undiluted mixture and incubated aerobically. When resistant strains of Bacteroides were mated with sensitive B. melaninogenicus, 0.05 ml of the undiluted, the  $10^{-3}$ , and the  $10^{-6}$  dilutions of the mixture were plated onto antibiotic-containing 5% blood agar. In all experiments involving B. melaninogenicus, the media was supplemented with a terminal concentration of 0.5  $\mu$ g of menadione per ml (Abbot Laboratories). B. melaninogenicus could be identified by pigment production. One multiply resistant strain, B. ochraceus 2228, was sensitive to  $2 \mu$ g of colistin per ml. This organism was mated with several sensitive strains of B. fragilis and suppressed from subcultures of the undiluted mixture by incorJ. BACTERIOL.

porating into the agar 10  $\mu$ g of colistin per ml, as well as the antibiotics under study.

Mating mixtures were plated onto 5% blood agar containing 10  $\mu$ g of either tetracycline, chloramphenicol, clindamycin, rifampin, or erythromycin per ml. In experiments with B. melaninogenicus, mixtures were also plated onto agar containing  $2 \mu$ g of penicillin per ml.

Bacteriocin experiments. Production of bacteriocins by two resistant strains of Bacteroides sp. were studied by the method of Fredericq (16). Prospective bacteriocinogenic strains were stabbed on plates of brain heart infusion agar and incubated for 48 h in a GasPak jar at 37 C. Enterobacter aerogenes K3151, a known colicin producer, was included on each plate. Colonies were sterilized over chloroform vapor, and a plasmid-negative strain of B. fragilis, B. melaninogenicus, or a colicin-sensitive E. coli K-12 was seeded over the surface of the plate from an overnight culture. The confluent cultures of the indicator strains were examined after 24 h of incubation for zones of inhibition surrounding the sterilized colonies.

#### RESULTS

Characteristics of Bacteroides strains tested. The 10 strains of Bacteroides selected for study included seven tetracycline-resistant B. fragilis (resistant to 20  $\mu$ g/ml), two tetracycline-sensitive B. fragilis (sensitive to 1.0  $\mu$ g/ ml), and <sup>1</sup> multiply resistant strain of B. ochraceus (syn. B. oralis subsp. elongatus) designated 2228. All strains were typically resistant to aminoglycoside antibiotics, polymyxins, and penicillin except B. ochraceus 2228, which was sensitive to  $2 \mu g$  of colistin per ml. All produced penicillinase. One strain of B. melaninogenicus was used as a sensitive recipient of drug resistance studies and was sensitive to 0.3  $\mu$ g of penicillin, tetracycline, chloramphenicol, clindamycin, rifampin, and erythromycin per ml. B. ochraceus 2228 was resistant to 20  $\mu$ g of tetracycline, chloramphenicol, clindamycin, erythromycin, and rifampin per ml. All strains of B. fragilis were sensitive to 1.0  $\mu$ g or less of these five drugs per ml except for tetracycline, as indicated above.

Of these 10 strains, two were shown to contain plasmids: a typical strain of B. fragilis subsp. fragilis (tetracycline resistant) designated N1175 and the multiply resistant B. ochraceus 2228. This strain of B. ochraceus was microaerophilic, producing visible colonies in a candle jar after 48 h of incubation. It did not produce spores, was consistently gram negative, and displayed the typical carbohydrate fermentation pattern of this organism (19), except that it fermented mannitol. In peptone-yeast-glucose broth it produced predominantly acetic, lactic, and succinic acids, as well as an unidentified short-chain alcohol with a retention time relative to acetic acid of 0.56.

The tetracycline-sensitive, plasmid-negative bacteria used in conjugation and bacteriocin experiments were designated B. fragilis A, B, and C.

Plasmids of B. fragilis strain N1175. The neutral sucrose gradient analysis of the cleared lysate from B. fragilis strain N1175 is shown in Fig. 1. Three discrete peaks of DNA, designated 1, 2, and 3, are seen. Using R100 and ColEl superhelical DNA as internal markers, the sedimentation coefficients of the three peaks were determined to be as follows: 1, 54S; 2, 41S; and 3, 22S. The S values of R100 and ColE1 are 77S and 23S, respectively, corresponding to molecular weights of  $70 \times 10^6$  and  $5 \times 10^6$  (2, 29). To assay for covalently closed circular DNA, the three peaks were pooled separately and sedimented through alkaline sucrose gradients (Fig. 2). Under these conditions, covalently closed circular DNA collapses into <sup>a</sup> rapidly sedimenting structure, whereas open circular and linear DNA denature into slower sedimenting single strands. Most of the DNA of peak <sup>1</sup> forms <sup>a</sup> sharp, rapidly sedimenting band characteristic of closed circular DNA (Fig. 2A), whereas the DNA in peaks <sup>2</sup> and <sup>3</sup> sediments much slower, approximately as expected for denatured, single strands (Fig. 2B and C). Analysis of peaks 1, 2, and 3 by CsCl-ethidium bromide equilibrium density gradient centrifugation confirmed that only peak <sup>1</sup> contained covalently closed circular DNA. The radioactivity in peak <sup>1</sup> represented between 1.0 to 1.2% of the total counts in the crude lysate.

The ratio of the sedimentation coefficients of



FIG. 1. Neutral sucrose gradient analysis of a cleared lysate from B. fragilis strain N1175. The lysate was prepared and sedimented through a 5 to 20% neutral sucrose gradient as described in Materials and Methods. "C-labeled covalently closed DNA forms of RIOO and ColEl were added as internal markers.



# FRACTION NUMBER

FIG. 2. Alkaline sucrose gradient analysis of the three DNA peaks shown in Fig. 1. Samples from each peak in Fig. <sup>1</sup> were layered directly onto separate 20 to 31% alkaline sucrose gradients and centrifuged as described in Materials and Methods. (A) Peak 1; (B) peak 2; (C) peak 3.

peaks <sup>1</sup> and 2 suggested that 2 is the open circular form of 1. To confirm this conclusion each DNA peak was examined by electron microscopy. The vast majority of the DNA from peak <sup>1</sup> appeared as individual superhelical molecules (Fig. 3A), as expected for closed circular DNA. Peak <sup>2</sup> consisted mostly of open circular DNA molecules (Fig. 3B) of <sup>a</sup> single size that corresponded to the open circular DNA form of the superhelical DNA in peak 1. In addition, a large number of heterogeneous linear DNA molecules were seen in peak 2, probably representing chromosomal DNA fragments. Peak <sup>3</sup> contained primarily linear DNA fragments but also a number of small open circular DNA molecules, No small superhelical DNA was seen, nor could any small covalently closed circular DNA molecules be isolated from CsClethidium bromide equilibrium density gradi-





FIG. 3. Electron micrographs of the B. fragilis plasmid DNA. (A) and (B) represent the DNA of peaks 1 and 2, respectively, described in Fig. 1. Both micrographs were taken at the same magnification.

ents carried out on cleared lysates. Thus, the strain of Bacteroides was studied because of its significance of these small DNA circles in peak multiple antibiotic resistance. Figure 4 shows significance of these small DNA circles in peak multiple antibiotic resistance. Figure 4 shows 3 is not clear.<br>  $\frac{1}{100}$  the CsCl-ethidium bromide density gradient of

the CsCl-ethidium bromide density gradient of Plasmids of B. ochraceus strain 2228. This the cleared lysate from strain 2228. The satellite



FIG. 3B

peak of closed circular DNA was pooled, dialyzed, and then centrifuged through a neutral sucrose gradient (Fig. 5). This gradient revealed two sharp plasmid peaks at 77S and 58S, with a broad peak of contaminating chromosomal DNA at <sup>25</sup> to 45S. Each plasmid peak contained about 1% of the total radioactivity present in the crude lysate. The superhelical nature of the plasmid peaks was confirmed by sedimenting the pooled samples from the gradients in Fig. 5 through separate alkaline sucrose gradients. The results (Fig. 6) demonstrate that



## FRACTION NUMBER

FIG. 4. CsCI-ethidium bromide equilibrium density gradient centrifugation of the cleared lysate from B. ochraceus strain 2228. The bar indicates the plasmid DNA fractions which were pooled for the neutral sucrose gradient analysis.



FIG. 5. Neutral sucrose gradient analysis of the plasmid DNA peak from Fig. 4. The pooled and dialyzed plasmid DNA was centrifuged through <sup>a</sup> neutral 5 to 20% sucrose gradient as described in Materials and Methods. "C-labeled superhelical  $R100$  and  $ColE1$  were added as internal markers.

the 77S and 58S peaks sediment rapidly as closed circular DNA species in alkaline sucrose, and that the 25 to 45S peak is probably contamination from the chromosomal band in the CsCI-ethidium bromide preparative gradient. The second rapidly sedimenting component in Fig. 6A represents contamination of the 77S peak with 58S DNA.

Conjugation experiments. The following attempts at transfer of antibiotic resistance were unsuccessful: (i) transfer of tetracycline resistance from B. fragilis N1175 to E. coli C, E. coli 53E, and Enterobacter cloacae 122H; (ii) transJ. BACTERIOL.

fer of tetracycline resistance and penicillinase production from  $B$ . fragilis N1175 to  $B$ . melaninogenicus; (iii) transfer of tetracycline, chloramphenicol, rifampin, erythromycin, and clindamycin resistance from B. ochraceus 2228 to B. melaninogenicus and B. fragilis A, B, and C; (iv) transfer of tetracycline and chloramphenicol resistance from B. ochraceus to E. coli C or E. coli 53E.

Bacteriocin production. Enterobacter aerogenes K3151 produced a bacteriocin (colicin V) active against  $E$ . coli C and  $E$ . coli K-12 but inactive against the plasmid-negative strains of B. fragilis and B. melaninogenicus. Sterilized colonies of B. fragilis N1175 and B. ochraceus 2228 did not inhibit either strain of Bacteroides,  $E.$  coli  $K-12$ , or  $E.$  coli  $C.$ 



FIG. 6. Alkaline sucrose gradient analysis of the three DNA peaks shown in Fig. 5. Samples from each peak in Fig. 5 were layered directly onto separate 20 to 31% alkaline sucrose gradients and centrifuged for 50 min at 50,000 rpm in the Spinco SW50.1 rotor at 15 C. (A) 77S peak; (B) 58S peak; (C) 35S peak.

# **DISCUSSION**

The demonstration of multiple plasmids from two strains of Bacteroides is of particular interest in view of the prominence of these bacteria among the normal flora. B. fragilis is by far the most common organism in the normal intestine (13, 14) and would be a logical reservoir for plasmids controlling the production of colicins, antibiotic resistance, or other properties vital to the maintenance of normal microbial ecology. There have been few studies of the distribution of B. ochraceus (B. oralis subsp. elongatus), but it appears to have been isolated primarily from subprimate animals (19).

B. fragilis strain N1175 contains a single species of covalently closed circular DNA with <sup>a</sup> sedimentation coefficient of 55S for the superhelical form, corresponding to a calculated molecular weight of  $25 \times 10^6$  (21). The plasmid DNA bands slightly less dense than the B. fragilis chromosome (40% guanine plus cytosine; J. Ryan, C. E. Davis, and A. I. Braude, unpublished data) in neutral CsCl equilibrium centrifugation. B. ochraceus 2228 was shown to contain two distinct closed circular DNA elements, the larger cosedimenting with the R100 marker, corresponding to a molecular weight of  $70 \times 10^{6}$  (25), and the smaller sedimenting at 58S, with a calculated molecular weight of 27  $\times$  $10<sup>6</sup>$  (21). Several cryptic plasmids recently isolated by Stiffler and colleagues from three strains of B. fragilis were much smaller, with molecular weights ranging from  $1.6 \times 10^6$  to  $4 \times$  $10<sup>6</sup>$  (30). Although we found no evidence of small closed circular DNA in our strains, it is possible that the open circular forms in peak 3 of B. fragilis N1175 represent closed circular DNA which has been converted to open circular DNA by the conditions of study. Similarly, low concentrations of small plasmids could be present in the 25 to 45S chromosomal peak of B. ochraceus 2228. In a recent abstract Damle and Syed (J. Dent. Res. 54:104, special issue A, 1975) reported 42 to 45S and 24 to 26S plasmid DNA in <sup>a</sup> strain of B. ochraceus and <sup>a</sup> <sup>24</sup> to 26S plasmid in B. melaninogenicus.

The physiological roles of these plasmids are unknown. Although the two plasmid-containing strains of Bacteroides were tetracycline resistant, we could not isolate closed circular DNA from six other resistant isolates. B. ochraceus 2228 was also resistant to chloramphenicol, erythromycin, clindamycin, and rifampin, drugs to which most Bacteroides are sensitive. Attempts to transfer antibiotic resistance from N1175 and 2228 to Enterobacteriaceae and other strains of sensitive Bacteroides were unsuccessful. Neither strain produced bacteriocins active against either a universally sensitive indicator strain of  $E$ . coli or plasmid-negative strains of Bacteroides. Although our experiments do not rule out the possibility that these plasmids are associated with resistance to antibiotics or other antibacterial agents, our inability to detect plasmids in six other strains of tetracycline-resistant B. fragilis with the method employed suggests that resistance to this antibiotic may not be plasmid-mediated in most strains of Bacteroides. These plasmids are large enough to contain sex factors, but it is possible that they mediate nontransmissible drug resistance.

Other possible functions of Bacteroides plasmids might include genetic control of properties important to bacteria that colonize mucosal surfaces. We plan to examine these strains for pili which could promote adherence to surfaces, for resistance to nitrites and other potentially toxic perservatives in food, and for resistance to heavy metals and the serum bactericidal system.

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