Synonymy of Strains of Center for Disease Control Group DF-1 with Species of Capnocytophaga

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Of eight strains of Center for Disease Control group DF-1 examined, seven had 62 to 87% deoxyribonucleic acid homology with the neotype strain of Capnocytophaga ochracea and one had 72% deoxyribonucleic acid homology with the type strain of C. gingivalis. Deoxyribonucleic acid homology of four strains of Bacteroides ochraceus with the neotype strain of C. ochracea was 76 to 86%.

In 1956 Prévot et al. (9) described two strains of obligately anaerobic, gram-negative rods which were morphologically similar to Fusiformis nucleatus (Knorr) Prévot (Fusobacterium nucleatum Knorr). These strains differed from the other strains of F. nucleatus examined in Prévot's study in that they produced ochre-redorange colonies in agar deeps, did not produce indole or a fetid odor, and were described as producing propionic, formic, and lactic acids. The strains were further characterized as fermenting carbohydrates to a low pH, reducing nitrates to nitrites, and not digesting gelatin, milk, or coagulated protein. The authors considered these two strains to be variants of F. nucleatus and named them F. nucleatus var. ochraceus. The strains, Pasteur Institute 1258B and 1411, had been isolated from purulent sputum and from a subcutaneous feline abscess caused by a bite from another cat, respectively.

In a later study from the Pasteur Institute, Sebald (11) examined the guanine plus cytosine content of many strains of anaerobic gram-negative rods maintained in the collection of the Service des Anaérobies and determined that F. nucleatus var. ochraceus (strain 1956C) had a guanine plus cytosine content of 42 mol% and that the guanine plus cytosine content of nine strains of F. nucleatus was 27.4 to 28.6 mol%. Because the base ratio of the strains of F. nucleatus var. ochraceus was similar to that which she found for species of Ristella (many members of which are now classified as Bacteroides in 'the 8th edition of Bergey's Manual [5]), Sebald proposed that F. nucleatus var. ochraceus be reclassified as Ristella ochracea.

B. oralis var. elongatus was described by Loesche et al. (6) in 1964 on the basis of a study of six strains isolated from the human oral cavity. Strains of B. oralis grew on the surface of

blood agar plates incubated in an anaerobic atmosphere but not on the surface of blood agar plates incubated in an aerobic atmosphere. B. oralis var. elongatus was differentiated from B. oralis var. oralis principally by cellular morphology. The authors further noted that the morphology of B. oralis var. elongatus was suggestive of fusobacteria and that some oral strains resembling B. oralis var. elongatus have "probably been characterized as Fusobacterium." The strains were described as fermenting carbohydrates and forming acetic and succinic acids. Strain R42 was designated the type strain.

The type strain of B. oralis var. elongatus, the two strains on which Prévot et al. (9) based the original description of F. nucleatus var. ochraceus, and strain 1956C used by Sebald (11) are no longer extant. One strain of B. oralis var. elongatus (SS31, received from Loesche) and three strains of R. ochracea (Prévot 2017B, 2476B, and 2376A) were compared at the Virginia Polytechnic Institute Anaerobe Laboratory. The strains all appeared to belong to the same species. The characteristics of these strains conformed well to the original descriptions of the respective varieties and, in the case of the strains from the Pasteur Institute, to the characteristics observed at the Pasteur Institute. All strains grew well and produced yellowish colonies on blood agar plates incubated in a candle extinction jar (3, 4). This was not considered to be inconsistent with the original descriptions, because no information was given in previous descriptions (6, 9) about the ability of strains to grow in conditions of reduced oxygen tension or increased carbon dioxide concentration. All four strains produced succinic and acetic acids from glucose, as had been reported by Loesche et al. (6). Since fermentation acids reported by Prévot et al. (9) probably were analyzed by Duclaux VOL. 10, 1979

distillation, which is sometimes difficult to interpret, the acetic acid could have been thought to be propionic and formic acids. None of the four strains reduced nitrates to nitrites, as had been reported originally for F. nucleatus var. ochraceus. However, copies of original data sheets listing reactions observed at the Pasteur Institute also showed nitrate-negative reactions for these strains, indicating that this reaction was considered to vary within the species. Because of the similarity among the strains, B. oralis var. elongatus and R. ochracea were considered subjective synonyms, and a new combination, Bacteroides ochraceus, was proposed (4, 8).

In the early 1960s Elizabeth 0. King, Center for Disease Control (CDC), differentiated a group of thin, gram-negative, saccharolytic rods which she designated the DF-1 group. Colonies of some strains were yellowish. On blood agar plates most strains grew better anaerobically or in the atmosphere of a candle extinction jar than they did in an aerobic atmosphere. The strains did not grow on MacConkey agar and were oxidase negative. They were described as fermentative, reducing nitrate, and not producing catalase or indole (12). The description of the DF-1 strains was similar to that of B. ochraceus (Table 1). The majority of the DF-1 strains were from human clinical specimens. It is highly probable that King did not recognize the close relationship between the DF-1 strains and B. oralis var. elongatus because the DF-1 strains were not obligately anaerobic, as B. oralis var. elongatus was presumed to be before 1972.

The genus *Capnocytophaga* has recently been proposed to include gliding, anaerobic to microaerophilic, gram-negative bacilli producing acetate and succinate as major metabolic end products and requiring carbon dioxide for both aerobic and anaerobic growth (S. C. Holt, E. R. Leadbetter, J. L. Simpson, E. D. Savitt, and S. S. Socransky, Arch. Microbiol., in press; E. R. Leadbetter, S. C. Holt, and S. S. Socransky, Arch. Microbiol., in press; S. S. Socransky, S. C. Holt, E. R. Leadbetter, A. C. R. Tanner, E. D. Savitt, and B. F. Hammond, Arch. Microbiol., in press; B. L. Williams and B. F. Hammond, Arch. Microbiol., in press). In previous deoxyribonucleic acid (DNA) homology studies by Williams and Hammond (in press), 27 of 28 strains of Capnocytophaga formed three mutually exclusive homology groups, represented by strains 25, 4, and 27. Because B. ochraceus ATCC ²⁷⁸⁷² and strain ²⁵ had DNA sequence homology at the level of 77%, they were considered to belong to the same species, and the new combination Capnocytophaga ochracea was proposed (Socransky et al., in press). New species C. sputigena and C. gingivalis were proposed for the genotypic and phenotypic groups represented by strains 4 and 27, respectively. In this communication we report the results of DNA homology studies with additional reference strains of B. ochraceus, representative strains from CDCdesignated DF-1, and the type or neotype strains of C. ochracea, C. sputigena, and C. gingivalis. Biochemical characteristics of the strains examined are also presented.

MATERIALS AND METHODS

Strains. Sources of strains used in this study are presented in Table 2.

DNA purification. Cultures for DNA purification were grown in Todd-Hewitt broth with 0.5% yeast extract. Cells from 1.5 liters of culture were harvested during late-log growth phase, washed with 0.05 M tris(hydroxymethyl)aminomethane, 0.05 M ethylenediaminetetraacetic acid, and 0.1 M NaCl (pH 8.0) and resuspended in the same buffer containing 25% (wt/ vol) sucrose. DNA was purified as described by Marmur (7) modified by the addition of 0.5 mg of pronase per ml (grade B, Calbiochem, San Diego, Calif.) during incubation with sodium lauryl sulfate. DNA was labeled in Capnocytophaga strains 25, 4, and 27 during growth in a medium containing amino acids, glucose, salts, and 5 μ Ci of [³H]thymidine (New England Nuclear Corp., Boston, Mass.). Labeled DNA was purified in the same manner as unlabeled DNA. The minimum specific activity achieved was $26,000$ cpm/ μ g (C. ochracea strain 25). Purified DNA was dialyzed against distilled water and sheared by ultrasonic treatment to an approximate molecular weight of 2.5×10^5 . All DNA preparations were stored frozen until used.

DNA hybridizations. DNA homo- and heteroduplex formations were assayed with the single-strand specific S1 endonuclease method of Crosa et al. (2). Hybridization mixtures of 150 μ g of unlabeled DNA and 0.1 μ g of ³H-labeled DNA were denatured at 100°C for ¹⁰ min in ¹ ml of 0.42 M NaCl. Reassociation conditions were 55°C for 16 h (T_m – 30°C for C. ochracea strain 25). Labeled DNA was similarly incubated in the absence of unlabeled DNA and with ^a known heterologous DNA (Escherichia coli K-12) to determine the levels of self-reassociation for each experiment. After incubation, duplicate portions (0.2 ml) were removed from the hybridization mixtures and subjected to 1,000 U of S1 endonuclease (Miles Laboratories, Inc., Elkhart, Ind.) for 20 min at 50°C. Duplicate portions were simultaneously incubated at 50°C but in the absence of the endonuclease. Radioactivity in double-stranded DNA segments was measured after termination of the S1 reaction and precipitation of DNA with cold 20% trichloroacetic acid (final concentration of 5%). The ratio of radioactivity in S1 treated and untreated portions was determined, and the results were normalized to the values of homologous reactions after correction for self-association of labeled DNA. Reported results are the average of at least three such hybridization experiments.

Physiological observations. Strains grown in bulk cultures for DNA purification were first tested

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for biochemical characteristics to ensure that DNA was purified from cultures of intended identities. Charlaboratory of B. L. W. Since the biochemical characteristics of individual reference strains are not availa-
ble in the literature, they are communicated here, essays and the laboratory of B. L. W. Since the biochemical characterization methods described are those used in the laboratory of B. L. W. Since the biochemical characterization methods described are those used in the lab

Figure 11. The interaction are considered and the extent of the interaction of the interaction methods described Preparation and inoculation of prereduced anaerobically sterilized (PRAS) basal peptone and yeast ex-COLOGO ...

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Preparation and inoculation of prereduced anaero-bically sterilize chemical tests were taken from 3-day PY cultures. A 4-ml amount of inoculum was suspended in 100 ml of freshly prepared (non-PRAS) PY basal medium of which ² ml was then aseptically dispensed into culture tubes previously autoclaved with 0.2 ml of $10\times$ concentrated substrate in distilled water; 5 ml of diluted inoculum also was added to 5 ml of $2\times$ gelatin medium (3). Final substrate concentrations were those listed in Figure 2.1 The substrate concentrations were those listed in the *Anaerobe Laboratory Manual* (3). Cultures were $\frac{3}{2}$ and $\frac{3}{2}$ the Anaerobe Laboratory Manual (3). Cultures were incubated in an anaerobic atmosphere of 10% hydrogen, 10% carbon dioxide, and 80% nitrogen achieved by evacuation and replacement. Acid production from carbohydrates was measured with a pH meter after 5 incubation at 35°C for 5 days. Assays for indole pro- $\frac{1}{2}$ $\frac{1}{2}$ were those described by Cowan (1). Metabolic end products were determined from PYG cultures as de scribed by Holdeman et al. (3) with the Hewlett-Packard (Avondale, Pa.) model 5830A gas chromatograph equipped with a flame ionization detector and a Supelco (Supelco, Inc., Bellefonte, Pa.) column of 15% $\frac{3}{5}$
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 copy of 24-h cultures in Todd-Hewitt broth (BBL Microbiology Systems, Cockeysville, Md.) supple mented with 0.5% yeast extract (Difco Laboratories, Detroit, Mich.). $\frac{1}{2}$
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EXECUTS

DNA sequence homology. DNA hybridization experiments with labeled DNA from the type of neotype strains of *C. cohracea, C. sputigena*, and *C. gingivalis* are summarized in Table 3. Eleven of the twelve *B. och* type of neotype strains of $C.$ ochracea, $C.$ spuexamples are summarized in the type of neotype strains of C. ochracea, C. spu-
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of the labeled DNA from C. ochracea. All four contracts and

two DF-1 strains showed more than 75% homology
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 $\frac{1}{2}$ at the intermediate level of 72%.
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Motility. All strains demonstrated gliding motility when log-phase cultures were observed in dark-field microscopy. The rate of transloca-

No. ^a	Designation	Source
VPI 5569	B. ochraceus	Prevot 2376C from actinomycosis (?) received as Ristella ochracea (Fusiformis nucleatus var. ochraceus).
VPI 5568	B. ochraceus	Prevot 2467B from dental suppuration, received as R. ochracea (F. nucleatus var. ochraceus).
VPI 5567	B. ochraceus	Prevot 2017B, received as R. ochracea (F. nucleatus var. ochraceus).
ATCC 27827	B. ochraceus	VPI 2845 from Loesche SS31 from human oral cavity (B. oralis var. elongatus).
CDC A7846	$DF-1$	Colorado State Health Department M418 from Mercy Hospital, Denver, CO, from throat.
CDC B2375	$DF-1$	Washington State Health Department Mc90 from Stevens Memorial Hospital, Edmons, WA, from human throat.
CDC B2906	$DF-1$	California State Health Department 5323-1-69 from Sequoia Hospital, Redwood City, CA, from blood.
CDC B5473	$DF-1$	W. J. Martin, Mayo University, Rochester, MN, C1293 from human blood.
CDC C4295	$DF-1$	Florida State Health Department CDC 426 from University Hospital, Jacksonville, FL, from finger.
CDC C6100	$DF-1$	Wisconsin State Health Department 63313 from St. Mary's Hospital, Madison, WI, from human vaginal specimen.
CDC C7282	$DF-1$	California State Health Department 2053 from human submaxillary gland.
CDC D6182	$DF-1$	Colorado State Health Department 76-1680 from L. Barth Reller, University of Colorado Medical Center, Denver, from human blood.
25	C. ochracea	S. S. Socransky, Forsyth Dental Center, Boston, MA, from juvenile periodontitis.
4	C. sputigena	S. S. Socransky, Forsyth Dental Center, Boston, MA, from rapidly advancing adult periodontitis.
27	C. gingivalis	S. S. Socransky, Forsyth Dental Center, Boston, MA, from juvenile periodontitis.

TABLE 2. Strains examined

" VPI, Virginia Polytechnic Institute and State Universitv; ATCC, American Type Culture Collection.

tion was 20 μ m per min for the most actively motile culture tested (DF-1 strain B5473).

Biochemical characteristics. The results of tests for carbohydrate utilization assayed by acid production, production of indole, and hydrolysis of gelatin are listed in Table 3. Acid production was scored as positive if cultures with the test substrate had ^a pH of at least 0.5 pH unit below the PY medium culture. All strains used in this

study produced acetic and succinic acids from PYG medium. Propionic and isovaleric acids were produced in low amounts (less than ¹ milliequivalent per liter) by some strains.

DISCUSSION

Anaerobic to microaerophilic, yellow- to orange-pigmented, gram-negative rods having features characteristic of organisms currently called

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B. ochraceus or CDC group DF-1 are being isolated with increasing frequency from clinical specimens, particularly blood cultures and oralfacial lesions. These show ^a high degree of DNA homology with species of the recently described genus Capnocytophaga. Of the 12 strains of B. ochraceus and CDC group DF-1 examined in this study, ¹¹ showed ^a high level of DNA homology with the neotype strain of C. ochracea, and one showed a high level of homology with the type strain of C. gingivalis.

Most strains showing significant homology to C. ochracea strain 25 also had a low level of sequence similarity with C. sputigena strain 4, as also has been observed with other strains of C. ochracea. The range of hybridization values of the DF-1 and B. ochraceus strains with C. ochracea 25 was similar to those observed among dental isolates of C. ochracea, but there was a proportionately larger number of DF-1 strains with lower hybridization values of 62 to 71% than had been observed with the dental strains. This is probably because the DF-1 strains chosen to include in this study reflected the extremes in phenotypic variation of strains that had been included in the group. This was done to try to establish the range of phenotypic variation that might occur among strains of the geno-species that would be determined by the homology studies.

The one DF-1 strain that showed a high level of DNA homology with C. gingivalis was ^a strain that fermented lactose, observed previously with only 8% of the strains studied. C. gingivalis strain D6182 also hydrolyzed starch and fermented galactose, which had not been reported previously for strains of this species (Socransky et al., in press).

None of the strains studied showed a high level of homology with C. sputigena. This is logical, assuming that the number of strains of these species reported by Socransky et al. (in press) is representative of the normal distribution of these species in the mouth. Of the strains studied by Socransky et al., only 10% (6 of 58) were C. sputigena. The distribution of the other two species, however, was nearly equal in their study, with 27 strains of C. ochracea and 25 strains of C. gingivalis.

The three species of Capnocytophaga are similar phenotypically. It appears that no single phenotypic test can be used to differentiate among them. In general, C. ochracea ferments the largest variety of carbohydrates tested and C. gingivalis the fewest (Socransky et al., in press). Phenotypic reactions of the three species have been reported previously (Socransky et al., in press).

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