Proposal for a Second Species Within the Genus *Ureaplasma*,
*Ureaplasma diversum* sp. nov.

C. J. HOWARD AND R. N. GOURLAY

A.R.C. Institute for Research on Animal Diseases, Compton, Near Newbury, Berkshire, England

It is proposed that a second species should be established within the genus *Ureaplasma*, *Ureaplasma diversum*, with strain A417 (NCTC 10182) as the type strain. The new species differs from *Ureaplasma urealyticum* serologically and on the basis of the guanine-plus-cytosine content of its deoxyribonucleic acid and its protein composition as determined by polyacrylamide gel electrophoresis. The species is serologically heterogeneous, the member strains apparently belonging to one of three serological clusters or groups.

The genus *Ureaplasma* comprises those members of the family *Mycoplasmataceae* that possess urease activity. There is a single species, *Ureaplasma urealyticum*, with strain T960 (ATCC 27618) as the type strain (35). In the definition of species within the genera *Mycoplasma* and *Acholeplasma*, considerable weight is placed on serological criteria (7). However, *U. urealyticum* was proposed as a serologically heterogeneous species, and in its original description, seven other strains, together with the type strain, were proposed as representatives of serovars (serotypes) I to VIII (35). Comparisons of these serovars by polyacrylamide gel electrophoresis (PAGE) has shown that many polypeptides are shared (34), whereas deoxyribonucleic acid (DNA) homology studies have shown a considerable homology between certain strains representing the serovars (4). These findings have confirmed the original proposal that *U. urealyticum* should be regarded as a single species with several serovars. Certain of these proposed serovars cross-react in some tests, whereas others appear totally distinct (1, 3, 6). The type strain of *U. urealyticum* and the strains of the other seven serovars were isolated from humans. Other isolates have been made from cattle and define their unique biological and serological properties. We hereby propose a second species within the genus *Ureaplasma*, with strain A417 (NCTC 10182) as the type strain.

**MATERIALS AND METHODS**

*Media.* Unless stated otherwise, isolates were grown in U4 broth or on U4 solid medium (19). The solid medium was modified slightly and contained 0.01% urea. Broth cultures were incubated aerobically at 37°C, whereas solid medium was incubated at 37°C under an atmosphere of 100% CO₂.

*Strains.* *U. urealyticum* strains 7, 23, 27, 58, 354, Pi, Co, and T960, serovars I to VIII (35), were obtained from F. T. Black, Aarhus, Denmark. The sources of bovine strains A417, Bu2, Mmb167, T95, T288, T45, D48, T315, T44, T71, and T74, considered to be representative of the serological diversity of bovine isolates, have been previously described (19); strain A417 originated from a pneumonic calf lung. All of these strains were cloned by filtering broth cultures (450-nm Millipore membrane) and picking single colonies at least four times.

Stock broth cultures of cloned strains were stored at −70°C and subcultured as required.

*Morphology.* Morphology was examined by using cells collected by centrifugation (10,000 × g for 30 min). Drops of suspensions on glass slides were air dried, fixed in methanol, stained with Gram or Giemsa stain, and examined by light microscopy.

For examination by electron microscopy (Phillips EM300; accelerating voltage of 80 kV), pelleted organisms were fixed with phosphate-buffered glutaralde-
Filtration studies. Numbers of color-changing units (CCU) in overnight cultures were determined in triplicate before and after passage of separate portions through membrane filters of 650-, 450-, 220-, and 100-nm average pore diameters (Millipore) at a positive pressure of 5 to 10 lb/in² (34.4 to 68.8 KN/m²).

PAGE. Details of the method used to examine [35S]methionine-labeled polypeptides from ureaplasma strains by sodium dodecyl sulfate-PAGE have been published (23).

Biochemical tests. The ability to catabolize urea or arginine was examined in broth cultures by inoculating isolates into U4 broth containing 0.05% urea, U4 broth with urea omitted and containing 0.5% arginine, or U4 broth with urea omitted and with no added arginine. The inoculated broths and uninoculated controls were incubated at 37°C and examined over 14 days for an alkaline color change indicating catabolism of either compound.

The ability to hydrolyze urea was also tested by a method similar to that of Swanberg et al. (38). Briefly, [14C]urea (57 μCi/μmol; Amersham, Great Britain) was added to U4 broth to give a final concentration of 1.25 μCi/ml. Pentuplicate inoculated broths and uninoculated controls were incubated at 37°C for 48 h. 0.1 ml of 3 N H2SO4 was added per ml of broth, and the broths were incubated in a fume hood at room temperature for 2 h, with occasional shaking, to remove CO2 (28). Samples (100 μl each) were mixed with 3 ml of scintillation fluid (KL402; Koch-Light Laboratories Ltd.), the radioactivity was counted in a Packard 2425 scintillation counter, and the percentage of urea catabolized was calculated.

Sterol requirement. The requirement for sterol was determined by a broth culture method similar to that described by Razin and Tully (30). Basal broth was prepared containing 70 ml of PPLO broth (Difco Laboratories, Detroit, Mich.), 10 ml of yeast extract (25% [wt/vol]), 0.25 ml of phenol red (1% [wt/vol]), 0.5 ml of thallium acetate (5% [wt/vol]), 0.5 ml of ampicillin (Beechams; 20% [wt/vol]), and 0.25 ml of urea (20% [wt/vol]). The following supplements were then added to give the indicated final concentrations: (i) fetal calf serum (20% [vol/vol]), (ii) bovine serum albumin (Armour fraction V; 0.5%) plus palmitic acid (10 μg/ml) together with either PPLO serum fraction (Difco; 1% [vol/vol]) or cholesterol at one of three concentrations (100, 25, or 5 μg/ml), (iii) bovine serum albumin plus palmitic acid without any cholesterol-containing supplement, and (iv) no added supplements. The volume was made up with water. Broths were inoculated with a washed suspension of ureaplasmata in 0.15 M phosphate-buffered saline (pH 7.2) containing 10² or 10³ CCU/ml and examined daily for growth, indicated by an alkaline color change in the medium.

A requirement for sterol was also determined in the indirect way by determining whether growth on agar was inhibited by disks containing 1.5% (wt/vol) digitoxin.

Serological tests. The metabolism inhibition (MI) test (14), growth inhibition (GI) test (19), and indirect immunofluorescent antibody (IF) test (18) were used with rabbit antisera. The IF test was used with antisera raised in gnotobiotic calves (16).
on solid medium, zones of inhibition of growth of strain A417 extended 7 mm from disks containing digitonin. No inhibition of growth was observed around disks not containing the compound. Similar results were observed with strains D48 and T44.

Strain A417 failed to grow in broth from which serum was omitted (basal broth). Furthermore, no growth was observed when bovine serum albumin and palmitic acid were added to the basal broth. Growth did occur in basal broth containing bovine serum albumin and palmitic acid plus PPLO serum fraction or cholesterol (at all three concentrations) and in basal broth containing serum.

**Biochemical characteristics.** Strain A417 produced a slight alkaline color change associated with a pH rise from pH 6.2 to pH 6.7 in U4 broth with urea omitted. In media made to the same formulation but containing 0.05% urea, a rise from pH 6.2 to 7.8 was produced, indicating that urea was catabolized. In broth not containing added urea but containing 0.5% arginine, the rise in pH after inoculation of strain A417 was similar to that in broth from which urea and arginine had been omitted. Thus, no evidence was seen for arginine catabolism.

In broth containing [¹⁴C]urea and inoculated with strain A417, <35% of the [¹⁴C]urea remained after incubation, compared with 100% in similarly incubated but uninoculated broths. This confirmed the presence of urease activity.

**Polyacrylamide gel patterns.** A comparison by PAGE of polypeptides from strains T960 and
VOL. 32, 1982

UREAPLASMA DIVERSUM

449

FIG. 3. Analysis of whole cell lysates of ureaplasma strains A417 and T960, grown in the presence of [35S]methionine, by sodium dodecyl sulfate-PAGE. Numbers across the top indicate strains. The two samples were run on the same gel. Arrows indicate dissimilar polypeptides.

A417 labeled by growth in the presence of [35S]methionine showed the two strains to be distinct (Fig. 3).

Previous studies had shown that serovars I to VII of U. urealyticum had many polypeptides common to strain T960, that representative bovine ureaplasmas had many polypeptides common to strain A417, and that the bovine isolates appeared distinct from the human ones (23).

DNA base composition. The G+C content of the DNA from strain A417 has been found to be 29.0 mol%. The range of values for 10 bovine isolates has been found to be from 28.7 to 30.2 mol% (20, 22).

Serological studies. Comparisons of bovine ureaplasmas by MI, GI, and IF tests with rabbit antisera indicated that three clusters of serologically similar, but not identical, strains existed (14, 16, 18, 29). Three strains, A417, D48, and T44, were selected as representatives of these three clusters, and it appears that bovine ureaplasmas can be identified by antisera raised in gnotobiotic calves to just three strains (16).

No serological relationship has been found between strain A417 and U. urealyticum strain T960 by the GI or IF tests with rabbit or calf antisera (14, 16, 18, 19, 29). Also, no relationship has been found between strain A417, or any other representative bovine ureaplasma isolate, and the eight serovars of U. urealyticum by these tests. A small, insignificant level of cross-reaction has been seen in the MI test between some bovine and human isolates (14, 41).

Habitat. Ureaplasmas representing one or more of the three serological clusters have been isolated from apparently normal and from diseased urogenital tracts from male and female cattle (33, 40, 42), from the eyes of cattle with keratoconjunctivitis (13), and from pneumatic calf lungs (10). The host range in other animals has not been determined.

Pathogenicity. Under controlled experimental conditions, strain A417 has been reported to produce mastitis in cows (12) and pneumonia in gnotobiotic calves (21). Other reports indicate that bovine ureaplasmas produce vulvitis (5) and conjunctivitis (32) experimentally in cattle. As inferred from ability or inability to produce mastitis experimentally, both virulent and avirulent strains exist in the cattle population (17). Thus, bovine ureaplasmas are potential pathogens.

Strain A417 has been shown to produce mastitis experimentally in goats (11); however, the relation between ovine and caprine ureaplasmas and bovine isolates remains to be elucidated. Ovine and caprine strains have been found to have G+C contents similar to, but perhaps marginally higher than, those of bovine isolates (22) but have been reported to be serologically distinct from bovine strains (25).

DISCUSSION

The results presented here indicate that strain A417 should be classified in the genus Urea-
plasma, but that this strain and other similar isolates from cattle are sufficiently distinct from the type strain of *U. urealyticum* and other similar human isolates to be regarded as a second species within the genus *Ureaplasma*.

The morphology of cells of strain A417 and the form of the colony, the absence of a cell wall, the resistance to penicillin and thallium acetate, the inability to grow at 25°C, the susceptibility to digitonin, the requirement for cholesterol, and the ability to pass through membrane filters indicate that strain A417 should be classified in the family *Mycoplasmataceae* (7). The possession of urease activity, the occurrence of optimum growth at pH 6.0 to 7.0, and the relatively small colony size indicate that this isolate should be classified in the genus *Ureaplasma* (35).

Since *U. urealyticum* is considered to be a species comprised of several serovars, it is clear that serological difference alone is not sufficient for proposing new isolates, from whatever source, as separate species. Nevertheless, it would seem to be necessary to show that a proposed second species was serologically distinct from the named species *U. urealyticum*. Strain T960, the type strain of *U. urealyticum*, is a member of serovar VIII. Other strains representing serovars I to VII were described at the same time as the type strain (35). Some cross-reactions occur between certain of the strains proposed as representative of serovars (1, 3, 6), and other strains have been isolated from humans that probably represent further serovars (26, 31).

Bovine ureaplasmas appear to exist as three serological clusters defined by rabbit antisera (18, 29) and as three serovars defined by gnotobiotic calf sera (16). The three clusters defined by rabbit antisera and the serovars defined by calf sera coincide.

Strain A417 is distinct from strain T960 and from members of other *U. urealyticum* serovars by a variety of serological tests, including MI, GI, and IF. In fact, no significant cross-reactions have been demonstrated among the eight *U. urealyticum* serovars and representative bovine isolates (16, 19, 29).

The G+C content of the DNA from *U. urealyticum* strain T960 was reported to be 27.1 mol% by Black et al. (2). Subsequently, the G+C content of strain A417 was reported to be 29.0 mol%. In the latter study, strain T960, which was included to make comparisons possible, was reported to have a G+C content of 27.4 mol% (20), a value virtually the same as that published previously. Although strains T960 and A417 have G+C contents that are different, the difference is not great. However, the range of values for eight *U. urealyticum* strains is 26.9 to 28.0 mol% (2), and this does not overlap the range (28.7 to 30.2 mol%) found for 10 bovine isolates (20, 22). These findings indicate that bovine isolates are a population distinct from those isolates classified as *U. urealyticum*.

Furthermore, the polypeptides of strains T960 and A417 are distinct when compared by PAGE. This method is considered to be suitable for distinguishing microorganisms at the level of species (7, 24). Other human isolates have been shown to have many polypeptides in common with strain T960, whereas other bovine isolates examined have many polypeptides in common with strain A417 but not with the human isolates (23, 34). Although some bovine isolates are distinct from strain A417 on the basis of serology, their polypeptides appear similar, and the bovine ureaplasmas are probably best regarded as belonging to one of three groups of the same species.

Thus, it is proposed that strain A417, which has been deposited in the National Collection of Type Cultures, London, England, as NCTC 10182, should be the type strain of the new species *Ureaplasma diversum* sp. nov. (L. adj. *diversum* different, distinct, heterogeneous; referring to the difference in polypeptides and G+C content, compared to *U. urealyticum*, and to the heterogeneous antigenic structure of the species).

As necessary (9), the proposal made here reflects the properties inherent in the strain and not the source of isolation; typically, however, *U. urealyticum* is not isolated from cattle and *U. diversum* is not isolated from humans.

As a consequence of these proposals, it would seem logical to compare isolates from other animal species with the representative *U. urealyticum* and *U. diversum* strains; as a minimum, antigenic structure, G+C content of the DNA, and the polypeptides should be examined. Although some strains are probably sufficiently similar to the two named species to be classified with them, some are almost certain to be sufficiently distinct to be regarded as further species.

ADDITIONUM

Since this manuscript was submitted, a study has been reported (C. Mouches, D. Taylor-Robinson, L. Stipkovits, and J. M. Bøve, Ann. Inst. Pasteur Paris 132B:171–196, 1981) which has confirmed by two-dimensional PAGE that bovine strains are distinct from those from humans and that they appear to fall into three groups coinciding with antigenic structure.

ACKNOWLEDGMENTS

We thank P. Bland for the electron micrograph, I. Jebbett for the photographs, and D. H. Pocock for his collaboration on the PAGE determinations.

REPRINT REQUESTS

Address reprint requests to: Dr. C. J. Howard, A. R. C.

LITERATURE CITED


