Mycoplasma imitans sp. nov. Is Related to Mycoplasma gallisepticum and Found in Birds

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A mycoplasma designated strain 4229T (T = type strain) was isolated in 1984 from the turbinate of a duck in France, and similar strains were isolated from geese in France and from a partridge in England. All of these strains were originally identified as Mycoplasma gallisepticum by immunofluorescence and growth inhibition tests, but subsequent serological and molecular studies indicated only a partial relationship to this species and DNA-DNA hybridization studies revealed only approximately 40 to 46% genetic homology with M. gallisepticum PG31T. In this study morphological, cultural, and physical investigations were carried out on strain 4229T and partridge strain B2/85, after we first demonstrated the similarity between these organisms by performing a restriction enzyme analysis of their DNAs. Both strains had phenotypic properties very similar to M. gallisepticum properties, including the presence of an attachment organelle. As a result of these investigations, the organisms were assigned to the class Mollicutes, the order Mycoplasmatales, and the genus Mycoplasma. They fermented glucose, reduced triphenyl tetrazolium chloride aerobically and anaerobically, and exhibited hemadsorption and hemagglutination, but other biochemical tests were negative. Apart from a serological cross-reaction with M. gallisepticum, these organisms exhibited no significant relationship with any previously described Mycoplasma species as determined by growth inhibition or immunofluorescence tests or with a number of additional serovars and unclassified avian strains. This Mycoplasma taxon therefore appears to be a new species, for which we propose the name Mycoplasma imitans. The type strain is strain 4229 (= NCTC 11733 = ATCC 51306). The significance of the organism has not been fully investigated, but preliminary in vitro and in vivo studies have indicated that it may be pathogenic.

Mycoplasma gallisepticum is a well-known respiratory pathogen of chickens and turkeys, and there have been occasional reports of isolation of this organism from other avian species, such as ducks (4, 27), geese (5), pheasants (32, 34), quails (3, 31, 34, 38), partridges (34, 42, 44), guinea fowl (40), pigeons (3), and peafowl (29).

In 1984 mycoplasmas were isolated from the turbinates of two mule (broiler) ducks in southwest France (15), and these mycoplasmas were identified as M. gallisepticum by immunofluorescence and growth inhibition (GI) tests. Later, the isolation of similar strains from geese in France was reported (11). Additional serological comparisons between these duck and goose isolates and some reference strains of M. gallisepticum in which GI and metabolism inhibition tests were used showed that the duck and goose strains were closely related to one another but were apparently less closely related to M. gallisepticum (16–18). Sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) profiles of the proteins, the results of a restriction enzyme analysis of the DNAs, and estimates of the G+C contents also suggested that the goose and duck strains were similar to one another but different from the reference strains of M. gallisepticum. Further support for these observations was provided by the results of Southern blot hybridization performed with rRNA and tuf gene probes (16, 45, 46), in which the goose and duck strains again appeared to be similar to one another but fell into a cluster separate from M. gallisepticum. Finally, it was shown by DNA-DNA hybridization that there was a very close genetic relationship between the duck and goose strains but that the levels of homology of these strains with M. gallisepticum were only 40 to 46% (17, 18).

Johnson (26) has suggested that taxonomic groups of organisms can be based on DNA homology data, and according to his recommendations, the duck and goose strains should not be regarded as members of M. gallisepticum, but as members of a closely related species. This is because the levels of homology are well below the proposed minimum value of 60%. The hypothesis that these strains should be placed in a separate species is strengthened by the proposal of an ad hoc committee of the International Committee on Systematic Bacteriology (41) that the phylogenetic definition of a species should be a taxon containing strains which exhibit approximately 70% or greater DNA-DNA relatedness.

Since strain 4229T (T = type strain) is distinct from M. gallisepticum, it may represent a new mollicute species, and in this study we used the methods recommended by the International Committee on Systematic Bacteriology Subcommittee on the Taxonomy of Mollicutes (25) to establish this. In most investigations we included another strain (B2/85) which we isolated from a partridge in England in 1985 and which we also originally identified by immunofluorescence tests as M. gallisepticum. We included this strain because it had also been used in previous investigations performed with the gene probes (45, 46) and appeared to be more closely related to the duck and goose strains than to typical M. gallisepticum.

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TABLE 1. Results of serological tests performed with Mycoplasma sp. strains 4229T and B2/85a

<table>
<thead>
<tr>
<th>Organism</th>
<th>GI test resultsb</th>
<th>Reference antiserum</th>
<th>Reference culture</th>
</tr>
</thead>
<tbody>
<tr>
<td>M. bovis Donetta</td>
<td>ND</td>
<td>0</td>
<td>ND</td>
</tr>
<tr>
<td>Mycoplasma bovoculi M165/69</td>
<td>ND</td>
<td>0</td>
<td>ND</td>
</tr>
<tr>
<td>M. clocosale 383</td>
<td>ND</td>
<td>0</td>
<td>ND</td>
</tr>
<tr>
<td>Mycoplasma columbinum MMPI</td>
<td>0</td>
<td>0</td>
<td>ND</td>
</tr>
<tr>
<td>M. dispar 462/2T</td>
<td>0</td>
<td>ND</td>
<td>ND</td>
</tr>
<tr>
<td>Mycoplasma fauca DC333T</td>
<td>ND</td>
<td>0</td>
<td>ND</td>
</tr>
<tr>
<td>M. gallisepticum PG31T</td>
<td>0</td>
<td>1(2)</td>
<td>1(1+)</td>
</tr>
<tr>
<td>Mycoplasma galiophyseum WR1</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>M. genitalium G3T</td>
<td>0</td>
<td>ND</td>
<td>s gl</td>
</tr>
<tr>
<td>M. h战胜uemoniae 1T</td>
<td>0</td>
<td>ND</td>
<td>g(w gl)</td>
</tr>
<tr>
<td>M. hyosynoviae S16T</td>
<td>0</td>
<td>ND</td>
<td>g(w gl)</td>
</tr>
<tr>
<td>M. lipojuc S171T</td>
<td>0</td>
<td>0</td>
<td>g(g)</td>
</tr>
<tr>
<td>M. lipophilum MaY</td>
<td>ND</td>
<td>0</td>
<td>ND</td>
</tr>
<tr>
<td>M. murs RIII</td>
<td>ND</td>
<td>0</td>
<td>ND</td>
</tr>
<tr>
<td>M. penetrans GTU-54-6A1</td>
<td>ND</td>
<td>0</td>
<td>ND</td>
</tr>
<tr>
<td>M. pneumoniae FH</td>
<td>ND</td>
<td>ND</td>
<td>ND</td>
</tr>
<tr>
<td>Mycoplasma pulmonis PG34T</td>
<td>ND</td>
<td>0</td>
<td>ND</td>
</tr>
<tr>
<td>Avian serovar J strain DIA</td>
<td>1(1)</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>Avian serovar K strain DK-CPA</td>
<td>0</td>
<td>0</td>
<td>g(g)</td>
</tr>
<tr>
<td>Avian serovar Q strain L3-10B</td>
<td>0</td>
<td>0</td>
<td>g(g)</td>
</tr>
<tr>
<td>Avian strain 700</td>
<td>0</td>
<td>ND</td>
<td>s gl</td>
</tr>
<tr>
<td>E. elychniae ELCN-1T</td>
<td>0</td>
<td>ND</td>
<td>ND</td>
</tr>
</tbody>
</table>


b The values indicate the sizes of inhibition zones (in millimeters).

0, negative result; wk, gi, weak glow; gl, glow; s, gl, strong glow; 1+, fluorescence. Reactions were considered positive only when fluorescence was observed.

ND, not done.

The data in parentheses are the results obtained with strain B2/85 or its antiserum. The other data are the results obtained with strain 4229T or its antiserum.

IFA tests performed with M. aortis, M. hyopharyngis, M. mobile, and M. pullorum also resulted in a glow with strain B2/85.

Antiserum prepared with strain RIII.
Pathology, Washington, D.C. *Mycoplasma lipophilum*, *Mycoplasma glycophilum*, and *Mycoplasma cloacale* reference strains were obtained from the collection of J. M. Bradbury. Avian strains 1220, 1223, and 19756 (members of three potential new species isolated from geese) and strain 700 (a member of a possible new species isolated from a chicken in Spain) were added to the collection of J. M. Bradbury by Z. Varga, Hungarian Academy of Science, Budapest.

**Restriction enzyme analysis.** In order to confirm that strains 4229T and B2/85 were genetically similar, restriction enzyme digests of the DNAs of these strains were compared. The DNA was extracted by a standard method (28), and the enzymes used were EcoRI, HindIII, and BglII.

**Media, cultivation, purification, and tests for reversion.** The media and conditions used to propagate most of the strains have been described previously (9). SP4 medium (39) was used for *M. leocapitus*, *M. muriis*, *M. penetrans*, *M. simbae*, *E. lict voxelax*, *E. luminosum*, *M. melaleucar*, and *E. somnilucar*, and *Mesoplasma lactucae*. Horse serum medium (13) was used to culture *Mycoplasma fermentans*, *Mycoplasma flocculare*, *Mycoplasma lipophilum*, *Mycoplasma mobile*, and *Mycoplasma pneumoniae*. Attempts to grow *E. elychniae*, *M. fastidiosum*, *M. genitalium*, *M. hyopneumoniae*, *Mycoplasma hyosynoviae*, *M. oxoniensis*, and *M. leopharyngis* were unsuccessful, and thus positive control cultures could not be included in the serological tests. Furthermore, *Mycoplasma dispar*, *Mycoplasma felinumutum*, and *M. spermatophilum* produced colonies that could be used in immuno-fluorescence tests, but we were unable to grow these organisms for GI tests.

*Mycoplasma* strains 4229T and B2/85 were subcultured routinely in broth or on agar (7) at 37°C in a carbon dioxide-rich (5%, vol/vol) atmosphere. They were purified by filter cloning them three times through filters with an average pore diameter of 450 nm (Nuclepore, Wallabs, Inc.). Tests for absence of reversion were carried out in broth without bacterial inhibitors. The culture of strain 4229T that was used for this experiment had been passaged seven times in our laboratory, but its passage history in France was not known. Strain B2/85 had undergone a total of seven passages in vitro. Ten passages of each strain were then made in broth without inhibitors, and each passage was examined for bacteria by subculturing it onto blood agar and by light microscopy.

**Morphology and filtration studies.** The colony morphology of strains 4229T and B2/85 was determined by light microscopy, and cell morphology was determined by light and electron microscopy (9, 12). Filterability studies were conducted by using 450- and 220-nm-pore-diameter membrane filters with starting cultures containing approximately 10^7 CFU/ml. The numbers of colony-forming units per milliliter remaining after filtration through membranes having different pore sizes were determined.

**Sterol requirement.** The effect of cholesterol on the growth of strain 4229T was investigated after we first adapted the organism to grow in serum-free broth containing 1% bovine serum fraction (33).

**Biochemical and biological properties.** Strains 4229T and B2/85 were examined for their susceptibility to digitonin and sodium polyanethol sulfonate (20, 22). The other biochemical tests performed included direct and indirect tests for utilization of glucose (1, 19) and tests for hydrolysis of urea (30, 37), hydrolysis of arginine, production of films and spots, phosphatase activity, tetrazolium reduction, liquefaction of coagulated horse serum (1), and hydrolysis of esculin and arbutin (23). Hemadsorption tests were conducted by using chicken, turkey, and duck erythrocytes, and hemagglutination activity was assessed by using chicken erythrocytes (24).

*M. gallisepticum* S6 was included in all of the tests described above as a positive or negative control culture, as appropriate, and in each test another reference species was selected as the other control. Thus, for example, in the test for hydrolysis of urea, *Ureaplasma gallorale* D6-1 was used as a positive control and *M. gallisepticum* was used as a negative control, while in the test for utilization of glucose, *M. gallisepticum* was the positive control and *Mycoplasma gallinarum* was the negative control.

**Serological tests.** Hyperimmune sera were prepared in rabbits against strains 4229T and B2/85 by using two rabbits for each strain. The method used was the method described previously (9), except that the initial numbers of intramuscular and subcutaneous inoculations were reduced from four to two and a booster intradermal inoculation containing 5 mg of antigen protein in Freund's incomplete adjuvant was given at 21 days. Other antisera were supplied in conjunction with the reference strains indicated above, and additional sera against several of the species were supplied by J. G. Tully.

The serological tests used were the indirect fluorescent antibody (IFA) test (35) and the agar well modification of the GI test (6). The antisera prepared in rabbits against strains 4229T and B2/85 were titrated by the IFA test; these antisera were subsequently used undiluted in GI tests and diluted 1/40 in IFA tests. One serum against each strain was also selected for cross-testing with *M. gallisepticum* PG31T and S6. A rabbit antiserum against 4229T prepared previously (18) was also included in these tests.

IFA and GI tests in which strains 4229T and B2/85 were compared with all of the previously described avian *Mycoplasma* species and serovars were performed in two ways: by using cultures of strains 4229T and B2/85 and the avian reference sera and by using all of the reference cultures with antisera prepared against strains 4229T and B2/85. These tests included five additional serovars of *Mycoplasma ioiae*, the three potential new species obtained from geese, and one species obtained from chickens.

To test strains 4229T and B2/85 with the previously recognized mammalian species, IFA tests were performed by using antisera against all of the mammalian reference strains. Positive controls were included except when we were not able or not permitted to grow the organism (e.g., *Mycoplasma mycoides* subsp. *mycoides*). For the GI tests cultures of the mammalian reference strains were tested with antisera against strains 4229T and B2/85 except for the species that we were not able to grow. For these species the reciprocal GI tests were performed by using reference antisera.

**RESULTS**

**Restriction enzyme analysis.** The DNA restriction patterns produced by strains 4229T and B2/85 were identical (Fig. 1) when all three enzymes were used, thus confirming the similarity of the two strains. These patterns were readily distinguishable from those of *M. gallisepticum* PG31T and S6.

**Growth characteristics.** Strains 4229T and B2/85 grew well in conventional mycoplasma broth and on agar. Strain 4229T grew more rapidly than strain B2/85 in broth medium. There was no evidence of reversion to bacteria after passage in broth lacking bacterial inhibitors.
FIG. 1. Restriction endonuclease cleavage patterns of strains 422T and B2/85 and *M. gallisepticum* PG31T and S6 digested with EcoRI (A), HindIII (B), and BglII (C). Lanes 1, phage lambda HindIII digest; lanes 2, PG31T; lanes 3, 422T; lanes 4, B2/85; lanes 5, S6. The sizes of the phage lambda molecular weight markers were as follows: a, 23.1 kbp; b, 9.4 kbp; c, 6.7 kbp; d, 4.4 kbp; e, 2.3 kbp; f, 2.0 kbp.

**Morphology and filtration studies.** Colonies having typical “fried-egg” morphology were produced on agar by both strains. Giemsa-stained films revealed coccoid forms, and phase-contrast examination also revealed mainly coccoid elements, but there was some evidence of pleomorphism. Electron microscopy of ultrathin sections revealed a three-layer membrane but no evidence of a cell wall (Fig. 2). The organisms were mainly oval or flask shaped, and a tiplike structure was clearly visible in both strains.

Both strains exhibited a decrease in concentration of 2 log10 CFU/ml after passage through a 450-nm-pore-diameter filter and a decrease in concentration of 3 log10 CFU/ml after filtration through a 220-nm-pore-diameter filter (Table 2).

**Sterol requirement.** As shown in Table 3, strain 422T exhibited a positive growth response to increasing amounts of cholesterol (strain B2/85 was not included in these tests). They were also susceptible to sodium polyethoxylate, but the inhibition zones were 2 to 3 mm smaller than the zones observed with digitonin. The organisms utilized glucose but gave negative results for hydrolysis of arginine and urea, production of films and spots, phosphatase activity, liquefaction of coagulated horse serum, and hydrolysis of esculin and arbutin. The mycoplasmas reduced tetrazolium both aerobically and anaerobically, but the reactions were weak. There was strong hemadsorption of chicken, turkey, and duck erythrocytes to colonies of strains 422T and B2/85, with approximately 75 to 100% of the individual colony surfaces covered. The *M. gallisepticum* control behaved similarly. In the hemagglutination test strains 422T and B2/85 and *M. gallisepticum* S6 gave reciprocal titers of 32, 16, and 32, respectively.

**Serological tests.*** Rabbit antisera prepared against strains 422T and B2/85 gave inhibition zones that were 5 to 7 mm wide in GI tests and reciprocal titers of 1,280 to 2,560 in IFA tests when they were tested with the homologous culture and with each other. IFA titers were considerably less when two strains of *M. gallisepticum* were used (Table 4).

The serological tests performed with antiserum to strain 422T and the heterologous avian *Mycoplasma* species and serovars gave negative results in GI and IFA tests except for the reactions shown in Table 1. The only results that were interpreted as positive were the immunofluorescence reactions with the type strain of *M. gallisepticum*. A similar reaction was observed in the reciprocal tests in which reference antiserum against *M. gallisepticum* was used, and a slight one-way reaction was observed in the GI test when the *M. gallisepticum* culture was used. The results obtained with strain B2/85 and its antiserum were virtually the same as the results obtained with 422T.

**DISCUSSION**

At the outset of this study a restriction enzyme analysis in which three different enzymes were used demonstrated that strains 422T and B2/85 were indistinguishable but were
FIG. 2. Electron micrographs of thin sections of *M. imitans* after 48 h in broth culture. (A and C) Strains 4229<sup>T</sup> and B2/85, respectively, showing round, elongated, and flask-shaped organisms. Bar = 1,000 nm. (B and D) Higher magnification of panels A and C, respectively, showing the three-layer membrane and terminal tip structure. Bar = 200 nm.

clearly different from two strains of *M. gallisepticum*. Previous studies (17, 18) had revealed differences in DNA restriction enzyme profiles between strain 4229<sup>T</sup> and reference strains of *M. gallisepticum*, but strain B2/85 had not been included in those comparisons.

The three enzymes were selected because they gave a large number of cuts in the DNA and might therefore be expected to reveal minor differences between strains. It could therefore be assumed that strains 4229<sup>T</sup> and B2/85 were closely related. This observation also substantiates the

<table>
<thead>
<tr>
<th>Strain</th>
<th>Before filtration</th>
<th>After filtration with 450-nm-pore-diameter filters</th>
<th>After filtration with 220-nm-pore-diameter filters</th>
</tr>
</thead>
<tbody>
<tr>
<td>4229&lt;sup&gt;T&lt;/sup&gt;</td>
<td>$7.2 \times 10^6$</td>
<td>$4.0 \times 10^6$</td>
<td>$1.6 \times 10^3$</td>
</tr>
<tr>
<td>B2/85</td>
<td>$6.7 \times 10^6$</td>
<td>$9.9 \times 10^4$</td>
<td>$3.6 \times 10^3$</td>
</tr>
</tbody>
</table>
TABLE 3. Growth response of strain 4229T to cholesterol

<table>
<thead>
<tr>
<th>Supplement(s) added to serum-free basal medium</th>
<th>Cholesterol concn (μg/ml) in medium</th>
<th>Protein yield (mg/100 ml)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Bovine serum fraction (control)</td>
<td>0</td>
<td>1.95</td>
</tr>
<tr>
<td>None</td>
<td>0</td>
<td>0.15</td>
</tr>
<tr>
<td>Bovine serum albumin</td>
<td>0</td>
<td>0.48</td>
</tr>
<tr>
<td>Bovine serum albumin, Tween, 80, and palmitic acid</td>
<td>0</td>
<td>0.43</td>
</tr>
<tr>
<td></td>
<td>1</td>
<td>0.81</td>
</tr>
<tr>
<td></td>
<td>5</td>
<td>1.08</td>
</tr>
<tr>
<td></td>
<td>10</td>
<td>1.20</td>
</tr>
<tr>
<td></td>
<td>20</td>
<td>1.40</td>
</tr>
</tbody>
</table>

data of Yogev et al. (45, 46) who examined these two strains with the pMC5 rRNA probe and the ruf gene probe and found that they produced similar genomic fingerprints.

Strains 4229T and B2/85 had properties consistent with membership in the class Mollicutes as defined by the International Committee on Systematic Bacteriology Subcommittee on the Taxonomy of Mollicutes (25). Thus, the colonies resembled fried eggs, and the cells were pleomorphic. There was no evidence of helical or motile forms and therefore no evidence that the organism belonged to the family Spiroplasmataceae. An electron microscope examination of thin sections revealed a three-layer membrane, and there was no evidence of a cell wall. The morphology of the organism was very similar to that of M. gallisepticum in that it possessed a tiplike structure. In other studies (36) we have shown that this structure has an attachment function.

The sterol requirement of strain 4229T was demonstrated by its growth response to cholesterol and was confirmed indirectly by the susceptibility of strains 4229T and B2/85 to digitonin. We therefore propose that this organism should be assigned to the family Mollicutes.

Since strain 4229T did not utilize urea, it is considered a member of the genus Mycoplasma rather than a member of the genus Ureaplasma, and the remaining biochemical and serological tests were carried out to provide a species description.

The G+C content of the DNA of strain 4229T has been estimated previously to be 31.9 mol% (17, 18); this value falls within the normal range of values for members of the Mollicutes but is lower by 1 to 2 mol% than the values for M. gallisepticum.

There was no serological evidence to suggest a relationship with any previously described Mycoplasma species other than the relationship already established with M. gallisepticum. Of the 107 mammalian and avian species, serovars, and strains compared by using GI tests with strains 4229T and B2/85, only M. gallisepticum and avian serovar J showed any sort of reaction, but zones were not considered large enough to be significant. In immunofluorescence tests, although several test results were recorded as glowing reactions, they were not considered positive, and the only positive reactions were the reactions seen with M. gallisepticum. Even these reactions were not as strong as had been originally recorded when the isolates were first (erroneously) identified.

Despite the many phenotypic similarities between strain 4229T and M. gallisepticum, the SDS-PAGE protein profiles and restriction enzyme analysis data for the DNAs of these organisms revealed obvious differences (18). It is interesting that Dox et al. (14), who developed a polymerase chain reaction for detection of M. gallisepticum, found that strain 4229T also gave a specific signal, although a commercially produced kit did not (8a). Thus, the choice of the DNA sequence used for amplification should be carefully assessed if distinction between these organisms is required. This finding is also supported by the results of another study (21) in which an oligonucleotide probe complementary to one of the variable regions in the 16S rRNA gene was used and the probe was found to react with strains 4229T and B2/85 and one of the goose strains.

We concluded that, despite the phenotypic similarities between strains 4229T and B2/85 and M. gallisepticum, the 40 to 46% level of genetic relationship between strain 4229T and M. gallisepticum shown by DNA-DNA hybridization studies cannot justified assigning these organisms to the same species (26). This situation is reminiscent of the situation which existed with Mycoplasma agalectae and Mycoplasma bovis. These organisms were originally assigned to the same species but were finally accepted as two distinct species on the basis of DNA-DNA hybridization values (approximately 40%) (2).

From these results we conclude that strain 4229T is a new Mycoplasma species, for which we propose the name Mycoplasma imitans. Strain B2/85, which was isolated from a different avian species and in a different country than strain 4229T, produced almost identical results in every test that was performed.

It is important to determine whether this new Mycoplasma species represents a threat to the poultry industry, either as a pathogen or as a complicating factor in serological testing programs for M. gallisepticum. Strains 4229T and B2/85 have been shown to cause ciliostasis in chicken and duck embryo tracheal organ cultures (36) and mortality of embryonated chicken, goose, and duck eggs (10, 16, 36). The results of preliminary in vivo studies have suggested that strain 4229T may cause reduced growth and airsacculitis in goslings and decreased liver size in older birds (10). Commercial turkeys experimentally infected by intranasal inoculation of strain 4229T when they were 2 days old showed sinusitis, corzya, conjunctivitis, and airsacculusitis when they were autopsied after 22 days (16). Mycoplasma-free turkeys infected with strain B2/85 by using eye drops and by inoculation into the thoracic airsac when they were 1 day old showed only transient mild respiratory signs (35a), but there was seroconversion and spread to contact birds. No clinical signs or lesions were observed in 8-week-old specific-pathogen-free chickens inoculated intratracheally with either strain 4229T or strain B2/85 or with one of the goose strains (43), but it is possible that the birds had developed natural resistance by this age. It is also possible that, like some strains of M. gallisepticum, the full pathogenic potential of the organisms may be realized only in the presence of other agents (8).

Description of Mycoplasma imitans sp. nov. Mycoplasma imitans (im'i.tans. L. v. gerund imitans, imitating, mimick-

TABLE 4. Cross-testing of strains 4229T and B2/85 and M. gallisepticum strains by immunofluorescence

<table>
<thead>
<tr>
<th>Strain</th>
<th>Reciprocal titers with the following antisera:</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>4229T</td>
</tr>
<tr>
<td>4229T</td>
<td>640</td>
</tr>
<tr>
<td>B2/85</td>
<td>640</td>
</tr>
<tr>
<td>PG31T</td>
<td>20</td>
</tr>
<tr>
<td>S6LP</td>
<td>40</td>
</tr>
</tbody>
</table>

* This antiserum had been prepared for a previous study (18).
ing, referring to the organism’s phenotypic resemblance to *M. gallisepticum*. Thin sections reveal oval and flask-shaped cells with a three-layer membrane and no outer wall. An attachment organelle is present. Cells pass through 450-nm-pore-size filters and through 220-nm-pore-size filters with some loss.

Typical “fried-egg” colonies are formed. The organism is resistant to penicillin and thallium acetate, and there is no reversion to bacterial forms in the absence of these compounds.

Cultures show a growth response to cholesterol, and growth is inhibited by digitonin.

Glucose is utilized, but arginine and urea are not utilized.

Triphenyl tetrazolium chloride is reduced aerobically and anaerobically.

There is no production of films and spots or phosphatase activity, no liquefaction of coagulated horse serum, and no hydrolysis of esculin or arbutin.

There is hemadsorption by chicken, turkey, and duck erythrocytes and hemagglutination of chicken erythrocytes.

The type strain is strain 4229 (= NCTC 11733 = ATCC 51306).

**ACKNOWLEDGMENTS**

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**REFERENCES**