

Pili (fimbriae) of *Branhamella* Species

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PURPOSE: Pili (fimbriae) have frequently been found to be involved in the attachment of bacteria to mucosal epithelial cells, an important initial step in the disease process. The purpose of this study was to determine if *Branhamella catarrhalis* expresses type 4 pili.

MATERIALS AND METHODS: Piliated *B. catarrhalis* phenotypic characteristics of colony morphology, agar corrosion, twitching motility, competence for deoxyribonucleic acid (DNA) transformation, autoagglutination, and pellicle formation were observed. DNA was isolated from *Branhamella* spp. and used in genomic Southern hybridizations with a *Moraxella bovis* pilin gene as a probe. Electron microscopy of negatively stained bacteria was carried out to visualize pili.

RESULTS: *B. catarrhalis* has several (but not all) of the phenotypic characteristics that are related to the presence of type 4 (MePhe) pili in closely related *Moraxella* spp., including competence for DNA transformation, autoagglutination, pellicle formation, colony morphology, and pitting of agar. The one phenotype we have not found that is generally characteristic of type 4 piliated bacteria is twitching motility. Genomic Southern hybridization analysis using a cloned *M. bovis* Q pilin gene as a probe reveals DNA homologous to the Q pilin gene in *B. catarrhalis*, *Branhamella ovis*, *Branhamella caviae*, and *Branhamella cuniculi*. Examination of *B. catarrhalis* strain ATCC25240 by electron microscopy reveals two different kinds of pili. One kind appears similar to other type 4 pili, whereas a second class is short pili extending outward from all portions of the bacteria.

CONCLUSION: Phenotypic, electron-microscopic, and hybridization data are all consistent with type 4 pili being present on some *B. catarrhalis* strains.

Bacterial colonization of mucosal surfaces depends upon attachment of bacteria to mucosal epithelial cells, often mediated by means of pili or fimbriae [1], which are protein filaments that extend from the surface of the bacteria. Pili are composed of up to 10,000 polymerized protein subunits called pilins. There are many different types of pili as defined by morphology and by functions such as pellicle formation in broth, and there are many types of pilins as defined by sequence comparisons [2]. In addition to pilin, some pili have one or more minor structural components [3,4], and in the case of the *Escherichia coli* P and type 1 pili, one of these minor subunits is the actual adhesin responsible for the binding of the pilus to the receptor molecule present on the eukaryotic cell surface [4-6]. Not only can a single bacterial species have strains that express structurally different classes of pili [2,7], but a single bacterium may have the capacity to express more than one class of pili, either sequentially or simultaneously [7,8].

Type 4 (MePhe) pili are present on a wide variety of pathogenic bacteria [9], including *Moraxella bovis* [10,11], *Moraxella nonliquefaciens* [12], *Neisseria gonorrhoeae* [13,14], *Neisseria meningitidis* [12], *Bacteroides nodosus* [15], *Pseudomonas aeruginosa* [16], and *Vibrio cholerae* [17]. As currently defined by *Bergey's Manual of Systematic Bacteriology*, the family *Neisseriaceae* contains the genera *Neisseria*, *Moraxella*, *Acinetobacter*, and *Kingella* [18]. A newly named genus, *Psychrobacter* [19], is also related to these genera. The genus *Moraxella* is subdivided into two subgenera, *Moraxella* and *Branhamella* [18]. *B. catarrhalis* was previously classed as *Neisseria catarrhalis* [18].

Our laboratory has been studying the type 4 (MePhe) class of pili present on *Moraxella* sp. and those of related genera. These studies include the cloning of the pilin gene from *M. bovis*, which causes bovine keratoconjunctivitis, and the use of this pilin gene to examine the presence and genetic structure of type IV pilin genes in other members of the *Neiseriaceae* [10] (Marrs CF, Stevens SP, Weir S, and Patel P, manuscript in preparation). This article discusses what we know about the pili of *Branhamella* spp.

TYPE 4 PILI IN MORAXELLA SPP.

The type 4 pilins all share extensive amino-terminal amino acid sequence homology and, with the exception of *V. cholerae*, they all contain the modified amino acid N-methylphenylalanine (MePhe) as the first residue of the mature protein [9,13,20]. That these conservations are important in a functional sense has been demonstrated by experiments in which *P. aeruginosa* bacteria processed and assembled *B. nodosus* pilins into pili structurally and immunologically indistinguishable from authentic *B. nodosus* pili [21,22]. Recently, similar experiments with our *M. bovis* pilin gene expressed in *P. aeruginosa* produced bacteria

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that actually had pili made up of both *M. bovis* pilin and *P. aeruginosa* pilin (Beard MKMG, Mattick JS, Moore LJ, *et al.*, manuscript submitted). Functionally, the pili of *M. bovis* attach to the corneal epithelium and only pilated strains are able to infect experimentally exposed cattle [11,23]. Similarly, pili appear to promote the pathogenicity of *N. gonorrhoeae* by mediating their attachment to human host mucosal surfaces [24].

MATERIALS AND METHODS

Bacterial Strains

B. catarrhalis strains ATCC8176, ATCC8193, ATCC25238, ATCC25239, and ATCC25240 were obtained from the American Type Culture Collection. *B. catarrhalis* strain B10 and *Moraxella lacunata* ATCC17956 were obtained from Dr. E. Juni, University of Michigan. *M. bovis* strain Epp63 was obtained from G.W. Pugh, Jr., Agricultural Research Service, Ames, Iowa. *N. meningitidis* strain M1080X was obtained from J.M. Koomey, University of Michigan.

Laboratory Techniques

Deoxyribonucleic acid (DNA) isolations and genomic Southern hybridizations were carried out as previously described [25]. All strains were grown on GC agar base (Difco Laboratories, Detroit, Michigan) with 1 percent IsoVitaleX (BBL Microbiology Systems, Cockeysville, Maryland). Two methods were used to assay for twitching motility, a hanging drop method of Depiazzi and Richards [26], and a plate method described by Henrichsen [27]. Electron microscopy of negatively stained bacteria was performed with a Zeiss EM-10CA transmission electron microscope using the procedures described by Hayat [28].

PHENOTYPES OF PILATED BACTERIA

For *N. gonorrhoeae*, *M. bovis*, *M. nonliquefaciens*, and other bacteria [9], type 4 piliation is associated with a variety of phenotypes. These include being located on the poles of the bacteria [9], colony morphology and agar corrosion [23,29], twitching motility [30,31], competence for DNA transformation [32], autoagglutination, and pellicle formation on the surface of broth cultures [33]. Of these phenotypes, autoagglutination and pellicle formation are also associated with other classes of pili [2], and bacteria like *Haemophilus influenzae* can be competent for DNA transformation in the absence of pili [34].

We have observed several of these phenotypes usually associated with type 4 pilated bacteria in our studies of *B. catarrhalis*. The colonies are similar in appearance to P⁺ *M. bovis* or *N. gonorrhoeae*. However, unlike *M. bovis* or *N. gonorrhoeae*, which show a readily detectable switch to a distinct colony type when they become non-piliated, we have yet to observe a *B. catarrhalis* with the normal nonpilated colony morphology. Because we have not yet isolated a *B. catarrhalis* that lacks pili, it is not clear how many of the following phenotypes we observe are specifically associated with being pilated. For all strains we have examined, *B. catarrhalis* colonies pit the agar, and the bacteria autoagglutinate and form pellicles in broth cultures. Most *B. catarrhalis* strains are capable of undergoing DNA transformation with related DNA [35,36] and in fact the ability of *B. catarrhalis* to be transformed by *M. nonliquefaciens* DNA was the

first evidence that it was related to *Moraxella* sp. [37]. Strain B10 is an exception, being non-competent for transformation despite forming a pellicle and having colonies that pit the agar and have P⁺ morphology. One characteristic type 4 piliation phenotype appears absent since two different twitching motility assays (described in Materials and Methods) were negative for every *B. catarrhalis* strain we tested, whereas the pilated controls from *M. bovis*, *M. lacunata*, and *N. meningitidis* all twitched. The presence of several phenotypes commonly correlated with type 4 piliation, coupled with the relatively close relationship between *B. catarrhalis* and several *Moraxella* spp. known to possess type 4 pili raised the question of whether *B. catarrhalis* might express type 4 pili.

GENETICS OF TYPE IV PILI

There are both similarities and differences in the genetic organization of type 4 pilin genes in different bacterial species. One similarity is that *P. aeruginosa*, *N. gonorrhoeae*, and *M. bovis* pilin genes all appear to use *rpoN* (σ^{54})-dependent promoters [38,39]. In terms of numbers of pilin genes, *P. aeruginosa* strains only have a single copy of the pilin gene in each genome [40,41], and most serotypes of *B. nodosus* also only have a single gene, but some are organized with the genes *fimA*, *fimC*, *fimZ* in order in the same orientation, where *fimA* and *fimZ* are both pilin genes [42]. *M. bovis* produces serologically different pilus types, and each strain is capable of producing one or the other of two pilus types [33,43]. *M. bovis* strain Epp63 expresses either Q pilin (formerly called beta) or I pilin (formerly called alpha). We have cloned and sequenced the *M. bovis* Q pilin gene [10], and shown that expression of either Q or I pilin alternates via a 2-kilobase inversion of genomic DNA [25]. *N. gonorrhoeae*, by contrast, contains multiple pilin gene loci in every strain and transitions from P⁺ to P⁻ and between different P⁺ pilin types are often accompanied by chromosomal DNA rearrangements [44-46]. Strain MS11 can express at least seven different pilin genes [45], has two regions of its chromosome that act as pilin expression loci [45], and has many other sites that contain silent variant pilin sequences [45,47]. Most strains each have a single pilin expression locus and multiple silent variant pilin sequences [48]. All silent copies are only partial genes, lacking the common N-terminal coding sequence of pilin [49,50]. Antigenic variation from one P⁺ type to another is a result of transformation of chromosomal DNA from other gonococci that have autolysed, followed by a *recA*⁺ dependent recombination event between silent copy donor DNA and the recipient expression locus [51]. Transitions from P⁺ to P⁻ sometimes involve deletions at the expression loci [45,49], and sometimes involve sequences from silent copy loci recombining into the expression locus to produce pilin molecules, which are defective in assembly [52].

As part of a general survey of bacteria related to *M. bovis* that we are carrying out, we hybridized our *M. bovis* Q pilin gene probe to genomic Southern blots containing DNAs from a variety of organisms. Figure 1 shows the hybridization patterns seen for four *Branhamella* spp. cleaved with *EcoRI*. Hybridization of the Q pilin probe to *Branhamella ovis* was moderately strong, about comparable with that seen with

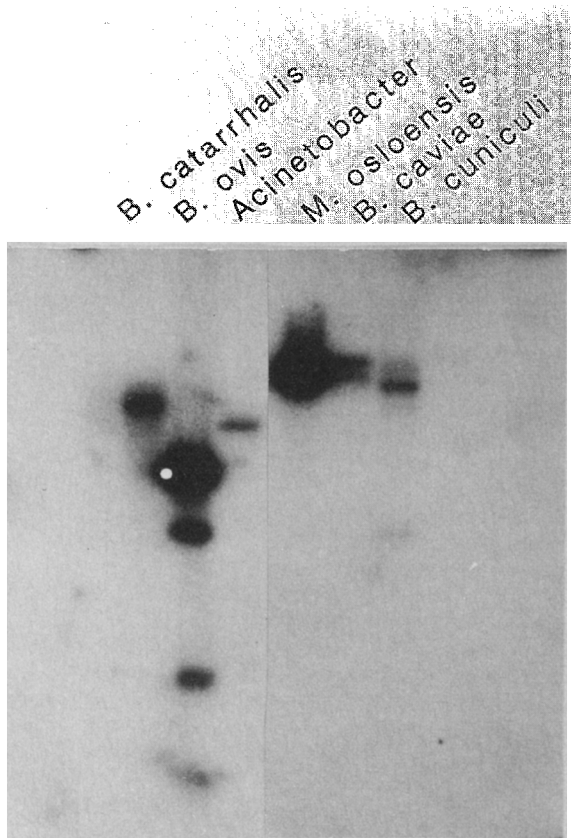


Figure 1. Hybridization using a *M. bovis* Q pilin gene probe to the following genomic DNAs cleaved with *Eco*RI: *B. catarrhalis* ATCC25238; *B. ovis* ATCC19575; *Acinetobacter calcoaceticus* ATCC15304; *M. osloensis* ATCC19962; *B. caviae* ATCC14659; *B. cuniculi* ATCC14688. Hybridization conditions were as described previously [44].

Moraxella osloensis (Figure 1), whereas *B. catarrhalis*, *Branhamella caviae*, and *Branhamella cuniculi* all hybridized at lesser, but clearly detectable levels. It is important to note that this hybridization data just tells us that DNA sequences similar to the *M. bovis* Q pilin gene are present in these bacteria. It does not tell us that type 4 pili are being expressed on the bacterial surface, or even guarantee that a complete pilin gene is present.

The observation that there are three bands of hybridization seen with *Eco*RI cleaved *B. ovis* DNA may mean that it contains more than one pilin gene. The fact that *B. catarrhalis* only has one band that hybridizes after *Eco*RI cleavage does not rule out it having more than one gene or partial gene. In *M. bovis* Epp63, the complete and partial pilin gene sequences are close enough that they appear on the same *Eco*RI fragment. It does, however, appear to rule out multiple, distinct pilin gene loci as is found in *N. gonorrhoeae* [48].

ELECTRON MICROSCOPIC EXAMINATION OF *B. CATARRHALIS*

In electron-microscopic studies, type 4 pili can be observed as long filaments extending from the pole of the cell [9,11]. An example of normal type 4 piliated bacteria is shown in Figure 2, top left.

The only report of pili being observed on *Branhamella* spp. was that of Wistreich and Baker [53], who reported that they failed to see any pili on two American Type Culture Collection strains of *N. catarrhalis* (ATCC8176, ATCC8193). In contrast, they reported that one *N. catarrhalis* strain from the culture collection of the University of Southern California had two different pili types. Some organisms had numerous short pili emerging from their outer borders, with an occasional long pilus extending beyond them, whereas with other bacteria the longer type predominated. Our observations of *B. catarrhalis* strain using the electron microscope match these descriptions quite well. Figure 2, top right and lower left and right, shows our electron micrographs of *B. catarrhalis* strain ATCC25240. In Figure 2, top right and lower left (at different magnification), can be seen examples of long, thin pili that appear very similar to the type 4 pili shown in Figure 2, top left. Figure 2, bottom right, shows the other class of short, thick pili with knobby ends that can be seen extending outward from all portions of the bacteria. Thus both phenotypic, electron-microscopic, and hybridization data all are in agreement that type 4 pili might be present on *B. catarrhalis*. The electron-microscopic data show that an additional, non-type 4 class of pili is also present. The relative roles of these two different classes of pili on the adherence properties and development of *B. catarrhalis* are exciting questions remaining to be explored.

REFERENCES

1. Beachey EH: Bacterial adherence: adhesin-receptor interactions mediating the attachment of bacteria to mucosal surfaces. *J Infect Dis* 1981; 143: 325-345.
2. Klemm P: Fimbrial adhesins of *Escherichia coli*. *Rev Infect Dis* 1985; 7: 321-340.
3. Oudega B, de Graaf M, de Boer L, et al: Detection and identification of FaeC as a minor component of K88 fibrillae of *Escherichia coli*. *Mol Microbiol* 1989; 3: 645-652.
4. Lindberg F, Lund B, Johansson L, Normark S: Localization of the receptor-binding protein adhesin at the tip of the bacterial pilus. *Nature* 1987; 328: 84-87.
5. Minion FC, Abraham SN, Beachey EH, Goguen JD: The genetic determinant of adhesive function in type 1 fimbriae of *Escherichia coli* is distinct from the gene encoding the fimbrial subunit. *J Bacteriol* 1986; 165: 1033-1036.
6. Maurer L, Orndorff PE: Identification and characterization of genes determining receptor binding and pilus length of *Escherichia coli* type 1 pili. *J Bacteriol* 1987; 169: 640-645.
7. Olsen A, Jonsson A, Normark S: Fibronectin binding mediated by a novel class of surface organelles on *Escherichia coli*. *Nature* 1989; 338: 652-655.
8. Archambaud M, Courcoux P, Labigne-Roussel A: Detection by molecular hybridization of *pap*, *afa*, and *sfa* adherence systems in *Escherichia coli* strains associated with urinary and enteral infections. *Ann Inst Pasteur/Microbiol* 1988; 139: 575-588.
9. Dalrymple B, Mattick JS: An analysis of the organization and evolution of type 4 fimbrial (MePhe) subunit proteins. *J Mol Evol* 1987; 25: 261-269.
10. Marrs CF, Schoolnik G, Koorney JM, Hardy J, Rothbard J, Falkow S: Cloning and sequencing of a *Moraxella bovis* pilin gene. *J Bacteriol* 1985; 163: 132-139.
11. Ruehl WW, Marrs CF, Fernandez R, Falkow S, Schoolnik GK: Purification, characterization, and pathogenicity of *Moraxella bovis* pili. *J Exp Med* 1988; 168: 983-1002.
12. Froholm LO, Sletten K: Purification and N-terminal sequence of a fimbrial protein from *Moraxella nonliquefaciens*. *FEBS Lett* 1977; 73: 29-32.
13. Hermodson MA, Chen KCS, Buchanan TM: *Neisseria* pili proteins: amino-terminal amino acid sequences and identification of an unusual amino acid. *Biochemistry* 1978; 17: 442-445.
14. Schoolnik GK, Fernandez R, Tai YT, Rothbard J, Gothschlich EC: Gonococcal pili: primary structure and receptor binding domain. *J Exp Med* 1984; 159: 1351-1370.
15. McKern NM, O'Donnell U, Inglis AS, Stewart DJ, Clark BL: Amino acid sequence of pilin from *Bacteroides nodosus* (strain 198), the causative organism of ovine footrot. *FEBS Lett* 1983; 164: 149-153.
16. Sastry PA, Pearlston JR, Smillie LB, Paranchych W: Amino acid sequence of pilin isolated from *Pseudomonas aeruginosa* PAK. *FEBS Lett* 1983; 151: 253-256.
17. Taylor RK, Miller VL, Furlong DB, Mekalanos JJ: Use of *phoA* gene fusions to identify a pilus colonization factor coordinately regulated with cholera toxin. *Proc Natl Acad Sci USA* 1987; 81: 2833-2837.
18. Bovre K: *Moraxella*. In: Kreig NR, Holt JG, eds. *Bergey's manual of systemic bacteriology*. Vol. 1. 1984; 296-303.
19. Juni E, Heym GA: *Psychrobacter immobilis* gen. nov., sp. nov.: genospecies composed of gram-negative, aerobic, oxidase-positive coccobacilli. *Int J Syst Bacteriol* 1986; 36: 388-391.
20. Elleman TC: Piliins of *Bacteroides nodosus*: molecular basis of serotypic variation and relationships to other bacterial piliins. *Microbiol Rev* 1988; 52: 233-247.

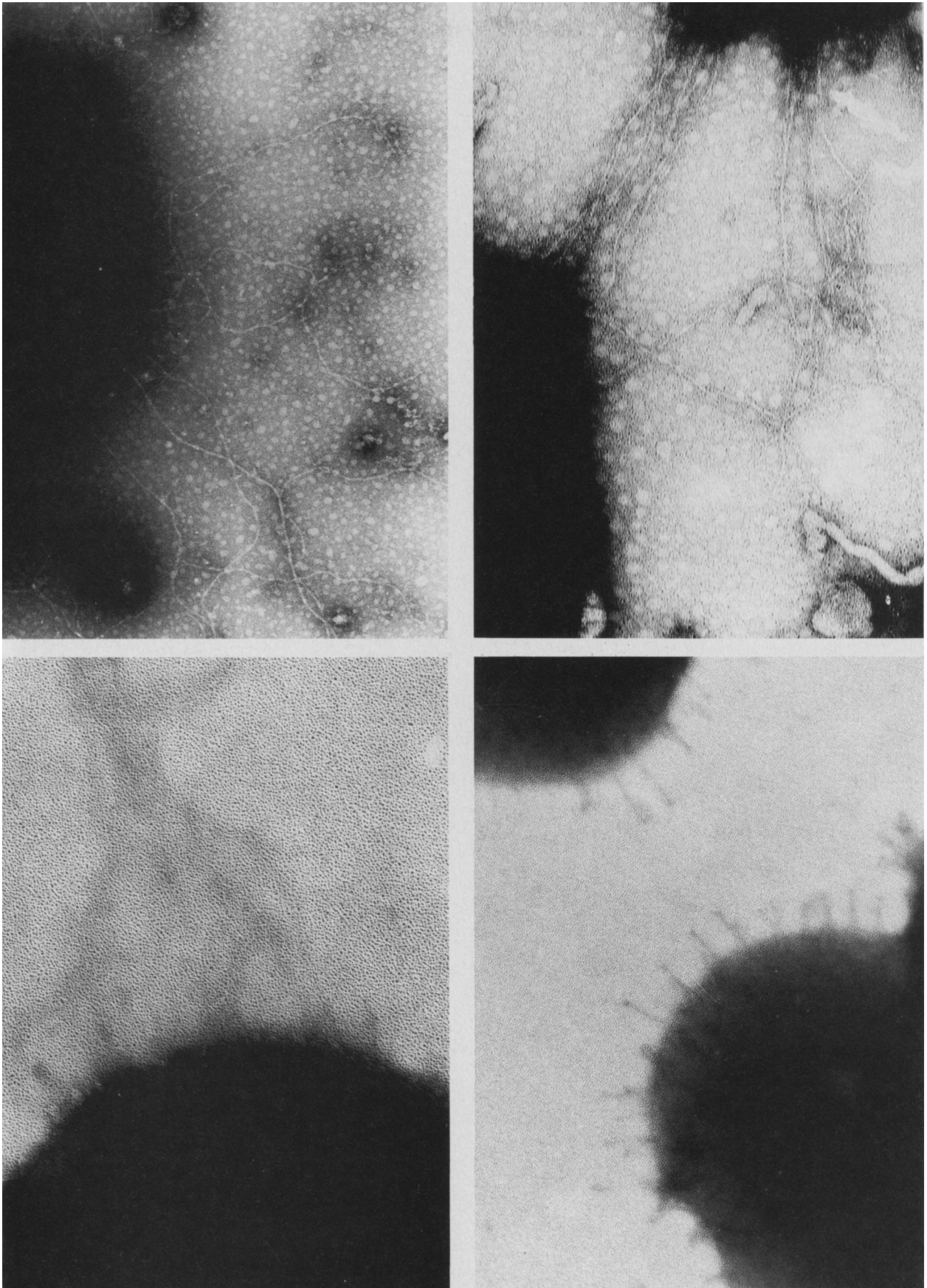


Figure 2. Top left, type 4 pili of *M. bovis* Epp63 P⁺, magnification $\times 94,000$; top right, long pili of *B. catarrhalis* ATCC25240, magnification $\times 140,000$; bottom left, potential type 4 pili of *B. catarrhalis* ATCC25240, magnification $\times 110,000$; bottom right, potential non-type 4 pili of *B. catarrhalis* ATCC25240, magnification $\times 91,500$.

21. Elleman TC, Hoynes PA, Stewart DJ, McKern NM, Peterson JE: Expression of pili from *Bacteroides nodosus* in *Pseudomonas aeruginosa*. J Bacteriol 1986; 168: 574-580.
22. Mattick JS, Bills MM, Anderson BJ, Dalrymple B, Mott MR, Egerton JR: Morphogenetic expression of *Bacteroides nodosus* fimbriae in *Pseudomonas aeruginosa*. J Bacteriol 1987; 169: 33-41.
23. Pedersen KB, Froholm LO, Borne K: Fimbriation and colony type of *Moraxella bovis* in relation to conjunctival colonization and development of keratoconjunctivitis in cattle. Acta Path Microbiol Scand [B] 1972; 80: 911-918.
24. Swanson J: Gonococcal adherence: selected topics. Rev Infect Dis 1983; 5 (suppl 4): S678-S684.
25. Marrs CF, Ruehl WW, Schoolnik GK, Falkow S: Pili gene phase variation of *Moraxella bovis* is caused by an inversion of the pilin genes. J Bacteriol 1988; 170: 3032-3039.
26. Depiazzi LJ, Richards RB: Motility in relation to virulence of *Bacteroides nodosus*. Vet Microbiol 1985; 10: 107-116.
27. Henriksen J: Bacterial surface translocation: a survey and a classification. Bacteriol Rev 1972; 36: 478-503.
28. Hayat MA: Basic techniques for transmission electron microscopy. New York: Academic Press, Inc., 1986; 232-234.
29. Bovre K, Froholm LO: Variation of colony morphology reflecting fimbriation in *Moraxella bovis* and two reference strains of *Moraxella nonliquefaciens*. Acta Path Microbiol Scand [B] 1972; 80: 629-640.
30. Henriksen J: Twitching motility. Ann Rev Microbiol 1983; 37: 81-93.
31. Henriksen J, Froholm LO, Bovre K: Studies on bacterial surface translocation. Acta Path Microbiol Scand [B] 1972; 80: 445-452.
32. Bovre K, Froholm LO: Competence in genetic transformation related to colony type and fimbriation in 3 species of *Moraxella*. Acta Path Microbiol Scand [B] 1972; 80: 649-659.
33. Sandhu TS, White FH, Simpson CF: Association of pili with rough colony type of *Moraxella bovis*. Am J Vet Res 1974; 35: 437-439.
34. Goodgal SH: DNA uptake in *Haemophilus* transformation. Ann Rev Genet 1982; 16: 169-192.
35. Bovre K: Affinities between *Moraxella* spp. and a strain of *Neisseria catarrhalis* as expressed by transformation. Acta Path Microbiol Scand 1963; 58: 528.
36. Catlin BW, Cunningham LS: Genetic transformation of *Neisseria catarrhalis* by deoxyribonucleate preparations having different average base compositions. J Gen Microbiol 1964; 37: 341-352.
37. Catlin BW: Reciprocal genetic transformation between *Neisseria catarrhalis* and *Moraxella nonliquefaciens*. J Gen Microbiol 1964; 37: 369-379.
38. Ishimoto K, Lory S: Formation of pilin in *Pseudomonas aeruginosa* requires the RpoN subunit of RNA polymerase. Proc Natl Acad Sci USA 1989; 86: 1954-1957.
39. Johnson K, Lory S: Characterization of *Pseudomonas aeruginosa* mutants with altered piliation. J Bacteriol 1987; 169: 5663-5667.
40. Pasloske BL, Finlay BB, Paranchych W: Cloning and sequencing of the *Pseudomonas aeruginosa* PAK pilin gene. FEBS Lett 1985; 183: 408-412.
41. Sastry PA, Finlay BB, Pasloske BL, Paranchych W, Pearlstone JR, Smillie LB: Comparative studies of the amino acid and nucleotide sequences of pilin derived from *Pseudomonas aeruginosa* PAK and PAO. J Bacteriol 1985; 164: 571-577.
42. Mattick JS: The molecular biology of the fimbriae (pili) of *Bacteroides nodosus* and the development of a recombinant-DNA-based vaccine. In: Egerton JR, Yang WK, Rifkin GG, eds. Footrot and foot abscess of ruminants, Boca Raton, Florida: CRC Press, 1989; 195-218.
43. Lepper AWD, Barton IJ: Infectious bovine keratoconjunctivitis: seasonal variation in culture, biochemical and immunoreactive properties of *Moraxella bovis* isolated from the eyes of cattle. Aust Vet J 1987; 64: 33-39.
44. Bergstrom S, Robbins K, Koomey JM, Swanson J: Piliation control mechanisms in *Neisseria gonorrhoeae*. Proc Natl Acad Sci USA 1986; 83: 3890-3894.
45. Hagblom P, Segal E, Billyard E, So M: Intragenic recombination leads to pilus antigenic variation in *Neisseria gonorrhoeae*. Nature 1985; 315: 156-158.
46. Meyer TF, Mlawer N, So M: Pilus expression in *Neisseria gonorrhoeae* involves chromosomal rearrangement. Cell 1982; 30: 45-52.
47. Segal E, Billyard E, So M, Storzbach S, Meyer TF: Role of chromosomal rearrangement in *N. gonorrhoeae* pilus phase variation. Cell 1985; 40: 293-300.
48. Swanson J, Bergstrom S, Barrera O, Robbins K, Corwin D: Pilus- gonococcal variants. J Exp Med 1985; 162: 729-744.
49. Haas R, Meyer TF: The repertoire of silent pilus genes in *Neisseria gonorrhoeae*: evidence for gene conversion. Cell 1986; 44: 107-115.
50. Segal E, Hagblom P, Seifert HS, So M: Antigenic variation of gonococcal pilus involves assembly of separated silent gene segments. Proc Natl Acad Sci USA 1986; 83: 2177-2181.
51. Seifert HS, So M: Genetic mechanisms of bacterial antigenic variation. Microbiol Rev 1988; 52: 327-336.
52. Swanson J, Bergstrom S, Robbins K, Barrera O, Corwin D, Koomey JM: Gene conversion involving the pilin structural gene correlates with pilus+ to pilus- changes in *Neisseria gonorrhoeae*. Cell 1986; 47: 267-276.
53. Wistreich GA, Baker RF: The presence of fimbriae (pili) in three species of *Neisseria*. J Gen Microbiol 1971; 65: 167-173.