JOURNAL OF BACTERIOLOGY, Sept. 1988, p. 3983–3990
0021-9193/88/093983-08$02.00/0

Vol. 170, No. 9

Gene Clusters for S Fimbrial Adhesin (*sfa*) and F1C Fimbriae (*foc*) of *Escherichia coli*: Comparative Aspects of Structure and Function

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Received 10 December 1987/Accepted 11 June 1988

Fimbrial adhesins enable bacteria to attach to eucaryotic cells. The genetic determinants for S fimbrial adhesins (*sfa*) and for F1C ("pseudotype I") fimbriae (*foc*) were compared. Sfa and F1C represent functionally distinct adhesins in their receptor specificities. Nevertheless, a high degree of homology between both determinants was found on the basis of DNA-DNA hybridizations. Characteristic differences in the restriction maps of the corresponding gene clusters, however, were visible in regions coding for the fimbrial subunits and for the S-specific adhesin. While a plasmid carrying the genetic determinant for F1C fimbriae was able to complement transposon-induced *sfa* mutants, a plasmid carrying the genetic determinant for a third adhesin type, termed P fimbriae, was unable to do so. Proximal *sfa*-specific sequences carrying the S fimbrial structural gene were fused to sequences representing the distal part of the *foc* gene cluster to form a hybrid cluster, and the *foc* proximal region coding for the structural protein was ligated to *sfa* distal sequences to form a second hybrid. Both hybrid clones produced intact fimbriae. Anti-F1C monoclonal antibodies (MAbs) only recognized clones which produced F1C fimbriae, and an anti-S adhesin MAb marked clones which expressed the S adhesin. However, one of four other anti-S fimbriae-specific MAbs reacted with both fimbrial structures, S and F1C, indicating a common epitope on both antigens. The results presented here support the view that *sfa* and *foc* determinants code for fimbriae that are similar in several aspects, while the P fimbriae are members of a more distantly related group.

Fimbriae or pili are bacterial cell wall appendages which consist of about 1,000 identical protein subunits. A single fimbria is 7 nm in diameter and up to 1 μm in length (9, 17). Very often fimbriae are associated with adhesins which enable the bacteria to attach to eucaryotic cells, including erythrocytes (RBC) (28, 31). Adherence of bacteria to eucaryotic cells is a prerequisite for many infectious diseases (27, 30, 41).

Extraintestinal *Escherichia coli* isolates cause infections of the urinary tract and the blood (sepsis) and are also the causative agents of newborn meningitis (32). Such *E. coli* isolates may carry different types of fimbrial adhesins. P fimbriae recognize an α-D-galactosyl-(1-4)-β-D-galactose receptor. They are strongly associated with uropathogenic *E. coli* strains and can be further subdivided into nine serologically distinct groups ($F7_1$ through F13) (16, 31, 41). The P fimbrial gene clusters of different serogroups are functionally related. This was shown by the construction of hybrid clones consisting of DNA sequences derived from different P determinants (*cis*-complementation) and also by the fact that P-specific gene products were able to produce a wild-type phenotype of P insertion mutants (*trans*-complementation) (24, 36, 43, 45). Another group of *E. coli* gene clusters, coding for type I fimbrial adhesins which interact with α-D-mannose-containing receptors, are also complementable by each other (5–7).

S fimbrial adhesins (Sfa) interact with α-sialyl-(2-3)-β-D-galactose-containing receptor molecules (21, 35). They are associated with urinary tract infection strains and, to a larger extent, newborn meningitis with *E. coli* isolates (22). The genetic determinant coding for Sfa has been cloned from the chromosome of a uropathogenic *E. coli* O6 strain (13). Recently, the S adhesin molecule, a protein of 12 kilodaltons (kDa), was isolated and characterized, and the sequence of the *sfaA* gene, coding for the S fimbrial protein subunit, was determined (28, 39). The *sfa* determinants from different strains have high sequence homology (33). Surprisingly, it was found that the *sfa* determinant is also related to another gene cluster (*foc*) coding for F1C fimbriae (34). In contrast to Sfa, however, F1C fimbriae are not able to agglutinate RBC (18, 44).

In this contribution, we present evidence that the *sfa* and *foc* determinants belong to the same general group of fimbrial gene clusters. The structural homology of the determinants and the serological relatedness of the corresponding proteins were demonstrated by the use of specific gene probes and monoclonal antibodies (MAbs). It is further shown that *foc*-specific gene products are able to produce a wild-type phenotype in *sfa* insertion mutants and that hybrid DNAs consisting of *sfa*- and *foc*-specific sequences code for intact fimbriae after transformation into nonfimbriated *E. coli* K-12 strains.

MATERIALS AND METHODS

Media, chemicals, and enzymes. Bacteria were grown in enriched nutrient broth or in alkaline broth extract. Radiochemicals were purchased from New England Nuclear Corp., Boston, Mass.; antibiotics were a gift from Bayer, Leverkusen, Federal Republic of Germany. All other chemicals were obtained from E. Merck AG, Darmstadt, Federal Republic of Germany. Restriction enzymes and T4 ligase were purchased from Bio-Rad Laboratories, Richmond, Calif. DNA polymerase I was obtained from Boehringer, Mannheim, Federal Republic of Germany.

Bacterial strains and recombinant DNAs. The wild-type strain 536 (O6:K15) exhibits the S fimbrial adhesin and type

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TABLE 1. Recombinant DNAs

Recombinant DNA	Genotype	Characteristics	Reference
pANN801-13	*sfa*$^+$ Ap^r	*sfa* determinant (4.1–13.9 kb)[a]	13
pANN801-11	Δ*sfa* Ap^r	Distal part of the *sfa* determinant (4.1–9.3 kb)	13
pANN801-15	Δ*sfa* Cm^r	Proximal part of the *sfa* determinant (11.1–13.7 kb)	33
pANN801-13-1	Δ*sfa* Ap^r	Proximal part of the *sfa* determinant (9.3–11.1 kb)	33
pMP13-Sph	Δ*sfa*	Proximal part of the *sfa* determinant (11.2–11.8 kb)	34
pANN1E6	*sfa* Ap^r	*sfa* determinant, *sfa*-flanking region, cosmid	Present study
pPIL110-54	*foc*$^+$ Cm^r	*foc* determinant (0.0–13.0 kb)	44
pPIL110-512	*foc*$^+$ Ap^r	*foc* determinant (5.9–7.8 kb)	44
pANN921	F8 Ap^r	F8 determinant	12
pRHU845	F13 (*pap*$^+$) Tc^r	F13 determinant	29
pANN801-C25	*foc sfa* Cm^r	Hybrid	Present study
pANN801-E4	*sfa foc* Ap^r	Hybrid	Present study

[a] The coordinates are given for *sfa* and *foc* determinants in Fig. 1.

I fimbriae and has been described elsewhere (11). The O18:K5 strain 2980 produces P fimbriae (serogroup F8) and F1C fimbriae; strain A21 (O7:K1) exhibits P fimbriae. The strains were described previously (15, 33, 46). For transformation and as controls in Southern hybridizations, the *E. coli* K-12 strains HB101 and 5K were used. The recombinant DNAs used in this study are listed in Table 1.

Recombinant DNA techniques. Chromosomal and plasmid DNA were isolated as described earlier (4, 10, 19). For restriction analysis, DNA was treated with appropriate restriction enzymes, and the resulting fragments were separated by gel electrophoresis on 0.7 to 1.0% agarose gels as described before (19). DNA fragments were isolated after agarose gel electrophoresis by electroelution (26). For cloning, DNA fragments were ligated into suitable vector molecules after heat inactivation of the restriction endonucleases at 65°C for 6 min as described before (3). Recombinant cosmid DNAs were selected from a recombinant cosmid gene bank of strain 536 by colony dot blot as described before (26). *E. coli* K-12 strains were transformed by the $CaCl_2$ method (23).

Generation of DNA probes. The *sfa*-specific gene probes are shown in Fig. 1. Probes A (coordinates, 0 to 4.4 kilobases [kb], Fig. 1), F (11.4 to 14.4 kb), and G (13.9 to 20.3 kb) were generated from the cosmid DNA pANN1E6. Probes B (4.1 to 7.1 kb) and C (7.1 to 9.3 kb) were derived from plasmid pANN801-11, probe D (9.3 to 11.1 kb) was from plasmid pANN801-13-1, and probe E (11.2 to 11.8 kb) was from plasmid pMP13-Sph (see Table 1). Only probes which were eluted from agarose gels were used in Southern hybridizations.

Nick translation, Southern hybridization, and autoradiography. The DNA probes were labeled by nick translation with a mixture of all four α-^{32}P-labeled deoxynucleoside triphosphates as described before (37). Transfer of DNA fragments from agarose gels to nitrocellulose filters and washing and autoradiography of the filters after hybridization were performed as described previously (40). The filters

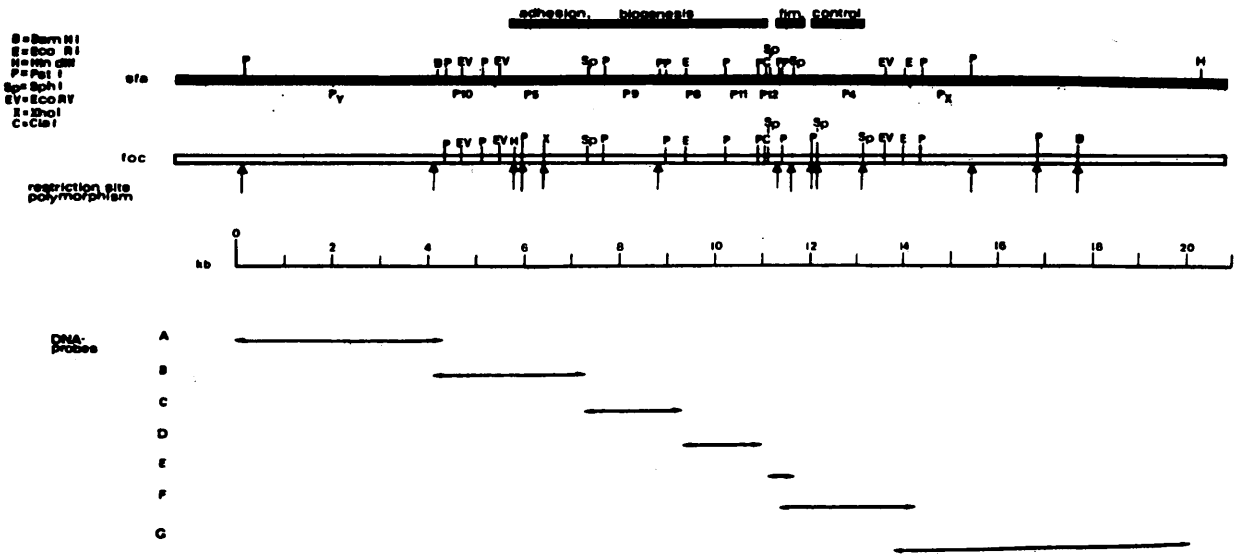

FIG. 1. Detailed restriction maps of *sfa* and *foc* determinants and their flanking regions. The *Pst*I fragments of the *sfa* determinants are indicated by numbers below the heavy line representing the *sfa* determinant. *Eco*RV, *Sph*I, *Cla*I, *Eco*RI, and *Xho*I sites are only shown for the coding regions. Differences between *sfa* and *foc* in restriction sites are indicated by arrows below the open bar representing the *foc* determinant. The DNA probes used in Southern hybridizations are indicated by the lines at the bottom (probes A to G). The functional regions of the *sfa* determinant are indicated at the top.

were hybridized in 50% formamide at 43°C for 3 days. Stringent conditions were used for the washing procedure: 30 min at room temperature in 2× SSC (1× SSC is 0.15 M NaCl plus 0.015 M sodium citrate)–0.1% sodium dodecyl sulfate (SDS) and then four times for 30 min each at 56°C in 0.1× SSC–0.1% SDS.

Preparation of fimbriae. Fimbriae were isolated from bacteria grown on plates as described previously (15).

SDS-polyacrylamide gel electrophoresis of fimbrial protein subunits. After fimbriae were disintegrated by boiling for 5 min in 10 mM Tris hydrochloride (pH 7.8) containing 4% SDS, 0.01 ml of mercaptoethanol per ml, 0.2 ml of glycerol per ml, and 0.002% bromphenol blue, the fimbrial samples (15 μg of protein) were run on slab gels as described previously (15, 46).

Preparation of polyclonal and monoclonal antisera. Polyclonal fimbria-specific antisera were prepared from rabbits with purified fimbrial protein eluted from an SDS-polyacrylamide electrophoresis gel (15). The preparation and characterization of the monoclonal antisera used were described elsewhere (1, 8, 28, 38).

Determination of fimbria antigens. Fimbriated cells were characterized by agglutination with antisera on glass slides. Western blots (immunoblots) were carried out by the method of Towbin et al. (42).

Immunoelectron microscopy. For immunoelectron microscopy, bacteria were transferred to copper grids coated with polyvinyl-Formvar. Incubation with MAbs and goat anti-mouse immunoglobulin G-coated gold spheres was described previously (28).

Determination of adhesins. S-specific adhesion was determined after mixing the bacterial cells with human and bovine RBC with and without 2% mannose and with RBC treated with neuraminidase (13, 21).

RESULTS

Genetic structure of *sfa* and *foc* determinants. The genetic determinants coding for S fimbrial adhesins (*sfa*) and F1C fimbriae (*foc*) were cloned on the recombinant plasmids pANN801-13 and pPIL110-54, respectively (13, 44). Recently it was shown by DNA-DNA dot blot hybridization studies that both gene clusters are structurally related (34). In order to demonstrate this relatedness more precisely, we generated five DNA probes (probes B to F, Fig. 1) which are specific for the *sfa* coding region. These probes were used in Southern hybridizations against *sfa*-, *foc*-, and P-specific recombinant DNAs (Fig. 2).

The *sfa* determinant was cleaved by the restriction enzyme *Pst*I into six fragments, P5, P9, P8, P11, P12, and P4. The main direction of transcription of the *sfa* determinant was from P4 to P5 (Fig. 1) (13). The *Pst*I-cleaved plasmid DNAs pANN801-13, pANN921 and pPIL110-54, coding for S fimbrial adhesins, P fimbriae, and F1C fimbriae, respectively, were hybridized against probe B, which represents the distal (3′) end of the *sfa* determinant. This region, in the case of *sfa*, codes for the S-specific adhesin (Fig. 1) (13, 28). It was demonstrated (Fig. 2) that the *Pst*I fragment P5 of *sfa* (lane A) disappeared in *foc* DNA, and two *foc*-specific fragments (ΔP5) were visible (lane B), indicating the presence of an additional *Pst*I site in the *foc* determinant compared with *sfa*. As summarized in Fig. 1, similar differences between *sfa* and *foc* determinants were found in that region on the basis of the sites for the restriction enzymes *Hin*dIII and *Xho*I.

After hybridization of *sfa*- and *foc*-specific DNAs with probes C, D, E, and F (Fig. 2 and data not shown), positive

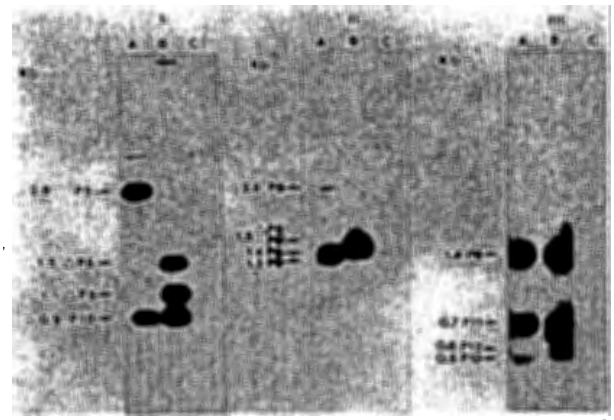

FIG. 2. Southern hybridization pattern of *Pst*I-cleaved DNA of plasmids pANN801-13 (*sfa*$^+$; lanes IA, IIA, and IIIA); pPIL110-54 (*foc*$^+$, lanes IB, IIB, and IIIB), and pANN921 (P-specific; F8, lanes IC, IIC, and IIIC; see Table 1). The DNAs were hybridized with nick-translated α-^{32}P-labeled DNA of probes B (blot I), C (blot II), and D (blot III) (see Fig. 1).

signals occurred in all cases tested. Only a few differences in restriction sites between *sfa*- and *foc*-specific DNAs (e.g., the absence of one *Pst*I site in *foc* compared with *sfa*) were visible in the regions corresponding to probes C and D, which code for proteins of 20, 26, and 90 kDa, involved in the biogenesis of the fimbriae (13; Ludwig and Hacker, unpublished results). The regions coding for fimbrial subunits (probe E and part of probe F) of both determinants, however, consisted of more heterogeneous stretches of DNA sequences, indicated in Fig. 1 (39, 44).

In contrast to the *foc* sequences, which hybridized to all *sfa*-specific probes used, the P-specific DNA showed no significant homologies with *sfa* probes B to E. However, homology was seen between P-specific sequences and *sfa* probe F (specific for the region involved in the control of transcription). A similar observation was also made recently by Ott et al. (34). These data correspond to the fact that about 1 kb of DNA of the control region of *sfa* and the P-specific determinant *pap* showed sequence homology of up to 90% in recent DNA sequence studies (2; T. Schmoll, W. Goebel, and J. Hacker, submitted for publication).

Genetic structure of *sfa* and *foc* flanking regions. In order to compare the DNA regions located adjacent to the *sfa* and *foc* gene clusters, an *sfa*-specific recombinant cosmid, pANN1E6, was isolated from a cosmid gene bank of strain 536 (20). pANN1E6 consists of the vector pHC79, the *sfa* determinant 6.5 kb in size, and DNA sequences flanking the *sfa* coding region at its proximal and distal ends. From these flanking DNA sequences, the DNA probes A (4.4 kb in size and specific for the 3′-flanking region) and G (9 kb in size and specific for the 5′-flanking region) were generated. These probes were hybridized against chromosomal DNAs isolated from strains producing F1C, P, or type I fimbriae.

The data presented in Fig. 3 clearly show that *Pst*I- (Fig. 3-I, lanes B and E) and *Bam*HI- (Fig. 3-II, lanes A and E) cleaved DNAs isolated from the F1C-positive strain 2980 and from the Sfa-producing wild-type strain 536 produced specific bands after hybridization against radioactively labeled probes A and G. In contrast, DNAs isolated from a P fimbria-specific strain (A21) and from two *E. coli* K-12 laboratory strains, which, in the case of strain 5K, exhibit type I fimbriae, gave no significant signals after being probed

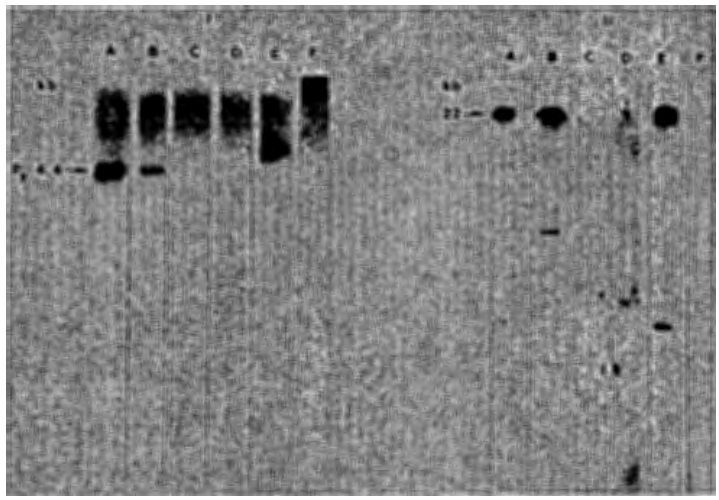

FIG. 3. Southern hybridization pattern of *Pst*I-cleaved (blot I) and *Bam*HI-cleaved (blot II) total DNA of strains HB101 (pcos1E6) (*sfa*$^+$; lanes IA and IIA), 536 (*sfa*$^+$; lanes IB and IIB), 5K (type I specific; lanes IC and IIC), A21 (P-specific; lanes ID and IID), 2980 (*foc*$^+$; P-specific; lanes IE and IIE), and HB101 (lanes IF and IIF). The DNAs were hybridized with nick-translated α-^{32}P-labeled DNA of probes A (blot I) and G (blot II); see Fig. 1.

with the DNAs representing *sfa*-specific flanking sequences. Therefore it can be concluded that not only the *sfa* and *foc* coding regions but also their flanking sequences, at least 12 kb (proximal) and 7 kb (distal to the coding regions) in size, showed strong homologies. These flanking sequences are not present on the chromosomes of strains carrying gene clusters coding for P and type I fimbriae.

***trans*-Complementation studies.** In order to find out whether the structural homology between *sfa* and *foc* gene clusters has consequences for their functional behavior, *trans*-complementation studies were carried out. In these experiments, P- and F1C-specific DNAs were tested for their ability to produce an Sfa$^+$ wild-type phenotype from Sfa$^-$ insertion mutants. Therefore, *E. coli* K-12 strains harboring transposon-induced mutations of the *sfa*-specific plasmid pANN801-13 were cotransformed with the *foc*-specific DNA pPIL110-54 and with the P-specific DNA pRHU845. The mutant DNAs carry Tn*1000* insertions in the *sfa* coding region of the *Pst*I fragments P9, P8, and P11 (Fig. 1). These sequences are responsible for the biogenesis of fimbrial adhesins (13, 34; Hacker and Ludwig, unpublished results). The different DNAs were able to coreplicate in the same cell because the pANN801-13 derivatives carry pBR322 sequences, whereas pPIL110-54 and pRHU845 consist of the vector molecule pACYC184 together with fimbria-specific insert DNAs (29, 44).

Table 2 summarizes the results of the experiments. The *sfa*-specific insertion mutants (pANN801-13/Tn*1000*-56, -57, -58) were complemented by the *foc*-specific plasmid pPIL110-54. This was shown by the expression of the S-specific fimbriae and an S-specific adhesion by the double transformants. The ability to express functions of the *sfa* determinant was restricted to the *foc*-specific sequences. Complementation was absent with cotransformants carrying the P-specific DNA pRHU845 and the *sfa* insertion mutants. Therefore it can be concluded that the *sfa* and *foc* but not P coding determinants are related at the functional level.

Construction of hybrid plasmids carrying *sfa* and *foc* coding sequences. In order to study further the functional relatedness of the *sfa* and *foc* determinants, hybrid plasmids containing DNA fragments derived from both gene clusters were constructed. The *sfa*-specific DNA pANN801-11, which carries a *Bam*HI-*Eco*RI fragment (coordinates 4.0 to 9.3 kb, Fig. 1) that includes the *Pst*I fragments ΔP8, P9, and P5, was cleaved with the enzyme *Eco*RI. pANN801-11 represents the distal (3′ end) region of the *sfa* determinant coding for the adhesive properties of the corresponding antigen. The *foc*-specific *Eco*RI fragment of pPIL110-512 (coordinates 9.3 to 13.9 kb) carrying a *focA* structural gene was ligated into pANN801-11, resulting in a hybrid plasmid, pANN801-E4. pANN801-E4 consists of the vector pBR322, the adherence-specific region of the *sfa* determinant including the gene *sfaS*, and the *foc*-specific sequences coding for the F1C fimbrillin protein FocA (Fig. 4).

In addition, the *sfa*-specific subcloned DNA pANN801-15, carrying the fimbrillin-coding gene *sfaA* of the *sfa* gene cluster on a *Cla*I-*Eco*RV fragment (coordinates 11.1 to 13.7 kb) inserted into pACYC184, was cleaved with the enzyme *Cla*I. A *foc*-specific *Cla*I fragment (coordinates ca. 0.0 to

TABLE 2. *trans*-Complementation between Tn*1000*-induced mutants of pANN801-13 (*sfa*$^+$), pPIL110-54 (*foc*$^+$), and pRHU845 (*pap*$^+$)

Recombinant DNA(s)	S fimbriae production[a]	S-specific adhesion[b]
pANN801-13	+	+
pANN801-13/Tn*1000*-56	−	−
pANN801-13/Tn*1000*-57	−	−
pANN801-13/Tn*1000*-58	−	−
pPIL110-54	−	−
pRHU845	−	−
pANN801-13/Tn*1000*-56, pPIL110-54	+	+
pANN801-13/Tn*1000*-57, pPIL110-54	+	+
pANN801-13/Tn*1000*-58, pPIL110-54	+	+
pANN801-13/Tn*1000*-56, pRHU845	−	−
pANN801-13/Tn*1000*-57, pRHU845	−	−
pANN801-13/Tn*1000*-58, pRHU845	−	−

[a] S fimbriae were detected by Western blots of fimbrial preparations isolated from *E. coli* clones blotted against monospecific anti-S polyclonal serum and by electron microscopy.

[b] Neuraminidase-susceptible hemagglutination of bovine RBC.

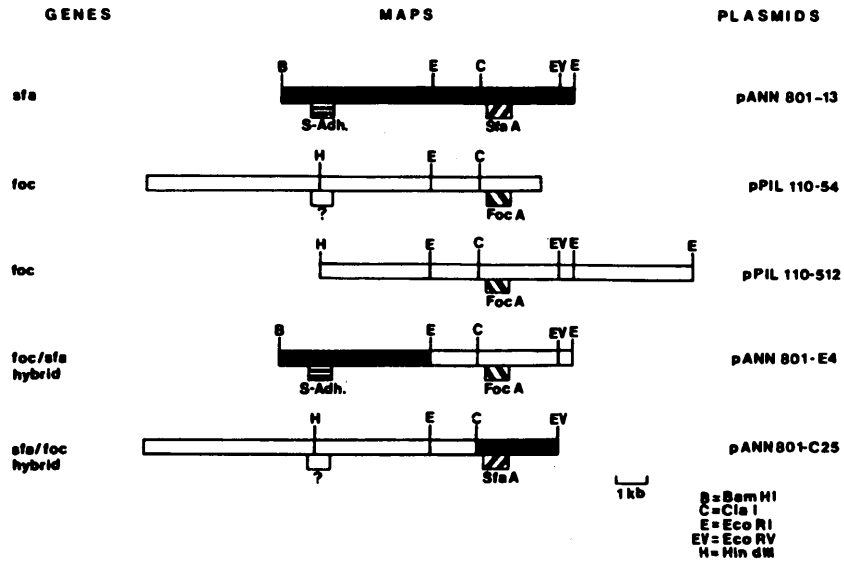

FIG. 4. Physical maps of the recombinant hybrid plasmids carrying *sfa*- and *foc*-specific DNA. The *sfa*-specific sequences are indicated by black bars, and the *foc*-specific DNA is marked by open bars. Fimbriae and adhesin genes are indicated by boxes below the bars. Only the cloning sites are shown. S-Adh, S adhesin.

11.1 kb) of plasmid pPIL110-54, representing the distal (3') part of the *foc* determinant, was ligated into the *Cla*I site of pANN801-15. The resulting hybrid plasmid, pANN801-C25, carries the fimbrillin-specific gene of *sfa* and the 3' end of *foc*. It was demonstrated by *Pst*I cleavage (data not shown) that the hybrid molecules carried the inserted DNAs in proper orientation (Fig. 4).

Characterization of the fimbrial clones. To answer the question of whether the recombinant clones produce intact fimbriae, the strains were analyzed by electron microscopy. As demonstrated in Fig. 5, the bacterial cells containing the parental *sfa*- and *foc*-specific DNA molecules or the hybrid plasmids were fimbriated. In order to identify the type of fimbriae produced, the cells were incubated with the F1C-specific MAb M9-6 (8) and then labeled with gold particles. As expected, the clones with a F1C-specific fimbrial subunit gene *focA* (pPIL110-54 and pANN801-E4) showed gold particles associated with the fimbriae, whereas the clones containing the S fimbrial structural gene *sfaA* did not. Identical results were obtained after using another F1C-specific MAb, F1-20025 (38) in Western blot experiments. Fimbriae prepared from the clones HB101(pPIL110-54) and HB101(pANN801-E4) reacted with the MAb in a protein band of 17 kDa (Fig. 6, blot II), which corresponds to the F1C protein subunit (44). In another Western blot (Fig. 6, blot I) analysis, the protein preparation of clones HB101(pANN801-13) and HB101(pANN801-E4) reacted with the S adhesin-specific MAb A1 in a band of 12 kDa. Therefore, one can conclude that the S-specific adhesin SfaS, which was described as a protein of 12 kDa very recently (28), is also produced by hybrid clones carrying the *focA* fimbrillin subunit gene.

In addition, four different antibodies (F1 to F4) which are specific for the native S fimbrial structure (28) were used to characterize the recombinant clones in agglutination tests. As shown in Table 3, three of these antibodies (F2 to F4) recognized clones HB101(pANN801-13) and HB101 (pANN801-C25). The recombinant DNAs of these clones carry the gene *sfaA*, coding for the S fimbrial subunit, a protein of 16 kDa (39). One MAb (F1), however, reacted with the S- and with the F1C-fimbriated clones, carrying the parental (pANN801-13, pPIL110-54) or the hybrid (pANN 801-E4, pANN801-C25) DNA. Thus, MAb F1 seems to recognize an epitope common for S and F1C fimbriae.

DISCUSSION

The two genetic determinants *sfa* and *foc*, coding for the S fimbrial adhesin and for F1C fimbriae, respectively, show similarities in their DNA sequence composition and exhibit common epitopes on their corresponding fimbrial proteins. In addition, *foc*- and *sfa*-specific DNA sequences were able to correct mutational defects of the corresponding gene cluster in *trans*- and *cis*-complementation experiments. Therefore it can be concluded that *sfa* and *foc* determinants form one main group of fimbrial determinants. In previous studies, two other general groups of fimbrial determinants were established on the basis of similar experiments: the gene clusters coding for P fimbriae and the determinants expressing type I fimbriae (5–7, 24, 36, 43, 45).

Sfa and F1C antigens, however, differ in their receptor specificities. While S fimbrial adhesins bind to the α-sialic acid-(2-3)-β-D-galactose residues of eucaryotic cell walls, F1C fimbriae do not bind to such substances (18, 21, 35, 44). On the other hand, it has been shown recently that F1C fimbriae are able to adhere to uroepithelial cells of human origin (Korhonen, personal communication). Therefore it is speculated that F1C fimbriae may recognize another receptor substance that is presumably slightly different from the S receptor molecule.

Very recently, a fimbrial determinant was cloned from the chromosome of an O75:K1 *E. coli* strain M. Pawelzik, J. Heesemann, J. Hacker, and W. Opferkuch, submitted for publication). This gene cluster, termed the "F1C-related fimbriae" (*sfr*) determinant, is also highly related to *sfa* and *foc* on the basis of DNA-DNA hybridizations and Western blots but is lacking the S-specific receptor-binding specificity. In addition, Lund et al. have described a fimbrial

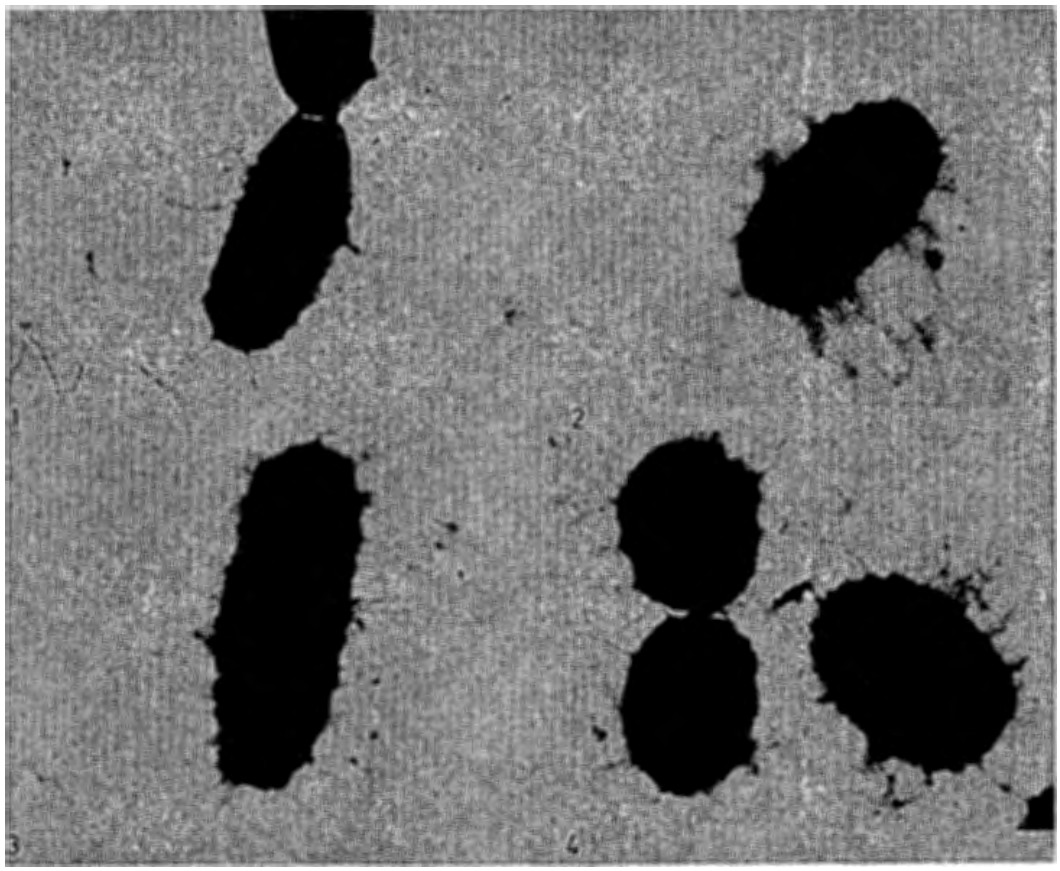

FIG. 5. Electron micrographs of *E. coli* K-12 strains carrying recombinant DNAs. The strains were incubated with F1C-specific MAb M9-6, followed by immunogold labeling and subsequent contrasting with 2% uranyl acetate. 1, HB101(pPIL110-54) (*foc*$^+$); 2, HB101(pANN801-13) (*sfa*$^+$); 3, HB101(pANN801-C25) (*sfa foc*); 4, HB101(pANN801-E4) (*foc sfa*).

determinant which is very similar to the *pap* gene cluster (P fimbriae of serogroup F13) in several aspects but codes for a different binding specificity (25). These data indicate that a shift in the receptor-binding specificity among one main group of fimbrial adhesins is a general phenomenon not restricted to Sfa and F1C fimbriae.

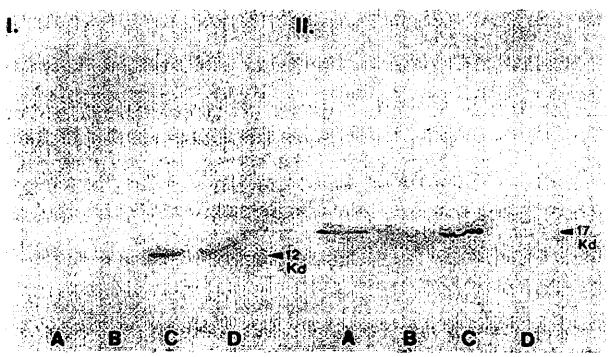

FIG. 6. Western blot of fimbrial protein preparations from strains HB101(pPIL110-54) (*foc*$^+$) (lanes A), HB101(pANN801-C25) (*sfa foc*) (lanes B), HB101(pANN801-E4) (*foc sfa*) (lanes C), and HB101(pANN801-13) (*sfa*$^+$) (lanes D). The fimbrial proteins were probed with the S adhesin-specific MAb A1 (blot I) and with the F1C-specific MAb F1-20025 (blot II). Kd, Kilodalton.

Differences between S fimbrial adhesins and F1C fimbriae were also established for the fimbrial subunit protein itself. The DNA sequences of the corresponding structural genes (*sfaA* and *focA*) show 73.3% homology (39, 44), but they differ in regions coding for the hydrophilic parts of the proteins, which should be responsible for their serological specificities (14). Therefore it was not surprising that five of six F1C or S fimbriae-specific MAbs only reacted with the homologous fimbriae (Table 3). One of these antibodies (MAb F1), however, recognized an epitope located on both fimbrial structures. This common epitope may be the reason for the serological cross-reactions between S and F1C fimbriae that were observed recently (33).

The antifimbriae MAbs reacted with the corresponding fimbrial subunit proteins isolated from *E. coli* K-12 strains carrying original recombinant plasmids or hybrid DNAs (Table 3, Fig. 5 and 6). These hybrid plasmids consist of DNA sequences specific for the *sfa* and *foc* gene clusters. It is interesting that an MAb which is specific for the S adhesin (MAb A1 [28]) also reacted with a hybrid clone that included a plasmid of *foc*-specific DNA which codes for the F1C fimbrial protein and a DNA region which harbors the S-specific adhesin gene (*sfaS*). These data show that the S

TABLE 3. Reaction of MAbs to recombinant *E. coli* strains expressing Sfa- or F1C-specific epitopes

Recombinant DNA	Fimbrial adhesin phenotype[a]	Reaction with MAb against:				
		S adhesin (A1)[b]	S fimbriae[c]		F1C fimbriae	
			F1	F2-4	M9-6[d]	F1-20025[b]
pANN801-13	Sfa-Fim$^+$/Sfa-Adh$^+$	+	+	+	−	−
pPIL110-54	F1C-Fim$^+$/F1C-Adh$^-$	−	+	−	+	+
pANN801-E4	F1C-Fim$^+$/Sfa-Adh$^+$	+	+	−	+	+
pANN801-C25	Sfa-Fim$^+$/F1C-Adh$^-$	−	+	+	−	−

[a] Fim, Production of fimbriae; Adh, adhesion to RBC; F1C fimbriae were set as Adh$^-$ (no agglutination of RBC [17, 44]).
[b] Western blot, see Fig. 6.
[c] Agglutination test.
[d] Electron micrograph after incubation of clones with immunogold-labeled antibodies; see Fig. 5.

adhesin molecules are coexpressed together with F1C fimbriae from the same hybrid gene cluster.

The functional relatedness of the *sfa* and *foc* gene clusters, demonstrated by the construction of hybrid DNA molecules (*cis*-complementations), was confirmed by *trans*-complementation studies. These experiments clearly show that *foc* gene products are able to express a wild-type phenotype, i.e., S-specific adherence, after cotransformation of *foc*-specific DNAs into strains which carry *sfa* plasmids with inserted Tn*1000* elements (Table 2). The fact that P-specific gene products did not express the wild-type phenotype of *sfa* insertion mutants is also supported by previous observations that P and *foc* sequences are unable to complement each other (36; Van Die, unpublished results).

It is remarkable that the homology between the *sfa* and *foc* determinants is not restricted to their coding regions. In Southern hybridization experiments, DNAs isolated from F1C- and Sfa-producing strains hybridized with gene probes specific for DNA sequences located in the vicinity but outside of the *sfa* gene cluster. From these data, it is obvious that the regions flanking the *sfa* and the *foc* determinant at both sides are also homologous to each other. It was further demonstrated by Southern hybridizations that these flanking regions (regions of about 20 kb were tested, Fig. 1 and 3) are specific for strains carrying *sfa* and *foc* determinants. These sequences are not present in the chromosomes of strains that produce P or type I fimbriae but lack Sfa or F1C fimbriae.

Similar results were observed for the *E. coli* hemolysin determinant (*hly*), a gene cluster coding for the α-hemolysin, another virulence factor of extraintestinal *E. coli* strains (27; J. Hacker, T. Schmoll, M. Ott, R. Marre, H. Hof, T. Jarchau, S. Knapp, I. Then, and W. Goebel, *in* E. Kass, ed., *Host-Parasite Interactions in Urinary Tract Infections*, in press). Hemolytic strains also carry *hly* determinants together with large flanking regions on their chromosomes. These tracts of DNA, which consist of the structural genes coding for α-hemolysin and about 70 to 100 kb of "flanking" sequences, are not present on the chromosomes of nonhemolytic strains. In addition, it was shown that the *hly* gene clusters and their flanking sequences located on the chromosomes of different *E. coli* strains exhibit a high degree of homology (20; Hacker et al., in press; Bender, Ott, and Hacker, unpublished results). Therefore it can be concluded that the occurrence and specificity of pathogenicity determinants together with large tracts of flanking sequences in the chromosomes of pathogenic *E. coli* strains are not restricted to the *foc* and *sfa* gene clusters.

ACKNOWLEDGMENTS

We thank Han deRee (Weesp, The Netherlands) for providing us with MAb M9-6, Staffan Normark (Umea, Sweden) for sending the plasmid pRHU845, and Bala Pillay (Würzburg) for critical reading of the manuscript. The excellent technical assistance of Heike Düvel is gratefully acknowledged.

The work was supported by the Deutsche Forschungsgemeinschaft (Ha1434/1-5).

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