Fimbriae- and flagella-mediated association with and invasion of cultured epithelial cells by *Salmonella enteritidis*

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***Salmonella enteritidis* expresses flagella and several finely regulated fimbriae, including SEF14, SEF17 and SEF21 (type 1). A panel of mutants was prepared in three strains of *S. enteritidis* to elucidate the role of these surface appendages in the association with and invasion of cultured epithelial cells. In all assays, the naturally occurring regulatory-defective strain 27655R associated with tissue culture cells significantly more than wild-type progenitor strains LA5 and S1400/94. Compared with wild-type strains, SEF14 mutants had no effect on association and invasion, whereas SEF17, SEF21 and aflagellate mutants showed significant reductions in both processes. Histological examination suggested a role for SEF17 in localized, aggregative adherence, which could be specifically blocked by anti-SEF17 sera and purified SEF17 fimbriae. SEF21-mediated association was neutralized by mannose and a specific monoclonal antibody, although to observe enhanced association it was necessary for the bacteria to be in fimbriate phase prior to infection. Additionally, aflagellate mutants associated and invaded less than motile bacteria. This study demonstrated the potential for multifactorial association and invasion of epithelial cells which involved SEF17 and SEF21 fimbriae, and flagella-mediated motility.**

Keywords: *Salmonella enteritidis*, fimbriae, flagella, tissue culture

INTRODUCTION

Salmonella enteritidis PT4 continues to be a major cause of foodborne infections in the United Kingdom and Europe, with poultry implicated as a major source (Humphrey & Rowe, 1988). Understanding the virulence mechanisms of bacterial pathogens depends upon the ability to study directly host–bacterial interactions. For *Escherichia coli* it is considered that adherence to enterocytes is a prerequisite for infection and the adhesive function of certain fimbriae has been described (Parry & Rooke, 1985). However, for *Salmonella* spp., the role of fimbriae and flagella is poorly understood, although it is commonly thought that they contribute to bacterial survival and persistence in the host and its environment (Thorns, 1995).

S. enteritidis has been demonstrated to express three morphologically distinct fimbriae and flagella. The

SEF14 fimbrial structure is <3 nm in diameter, is composed of repeating fimbrin subunits 14·3 kDa in size and is unique to certain serotypes in serogroup D, including all strains of *S. enteritidis* and *Salmonella dublin* (Thorns *et al.*, 1990, 1992). SEF17 are hydrophobic organelles (Collinson *et al.*, 1991) that mediate binding to the tissue matrix fibronectin (Collinson *et al.*, 1993; Baloda *et al.*, 1988), plasminogen (Sjöbring *et al.*, 1994) and human-contact-phase proteins (Ben-Nasr *et al.*, 1996; Herwald *et al.*, 1998). Type 1 fimbriae are ubiquitous amongst members of the *Enterobacteriaceae* and mediate mannose-sensitive haemagglutination (MSHA) (Duguid *et al.*, 1966; Müller *et al.*, 1991) with animal erythrocytes. Type 1 fimbriae of *S. enteritidis* (SEF21) and *Salmonella typhimurium* have been reported recently to bind to rat gut epithelium (Ewen *et al.*, 1997).

The majority of *Enterobacteriaceae* elaborate flagella, which are considered to assist the bacterium with directional motility, although their role in the pathogenesis of *Salmonella* is equivocal. A mutant of *S.*

Abbreviation: MSHA, mannose-sensitive haemagglutination.

typhimurium deficient in both MSHA and motility was reported to be greatly reduced in its ability to invade epithelial cells *in vitro* and persist in the liver and spleen of orally challenged chicks, whereas mutants defective for the elaboration of either flagella or fimbriae alone showed no significant effect (Lee *et al.*, 1996).

Additionally, *S. enteritidis* has the genetic potential to elaborate a plasmid-encoded fimbriae (PEF) analogue, harboured by the *S. enteritidis*-associated virulence plasmid (Baumler *et al.*, 1996a) and long polar fimbriae (Baumler & Heffron 1995; Baumler *et al.* 1996b). The structure of long polar fimbriae has not been described in *S. enteritidis*, whereas a serological response to PEF has been reported *in vivo* (Woodward *et al.*, 1996), although plasmid-free organisms remain fully virulent in chickens (Halavatkar & Barrow, 1993). *Salmonella* fimbriae are finely regulated and expression can be modulated *in vitro* by varying temperature, pH, surface contact (Dibb-Fuller *et al.*, 1997; McDermid *et al.*, 1996; Walker *et al.*, 1999) and available nutrients (Thorns *et al.*, 1992).

The interaction between pathogen and host cells during the onset of disease is a current focus of our research. Specifically, very little is known about the contribution of surface appendages of *S. enteritidis* in colonization of the chick. This report describes, as part of these studies, the use of defined mutants to investigate the individual and collective role of fimbriae and flagella in adherence and invasion of cultured epithelial cells prior to *in vivo* studies in poultry.

METHODS

Bacteria. *S. enteritidis* strains S1400/94 and LA5 are wild-type strains isolated from natural chicken infections and were obtained from a reference collection at the Veterinary Laboratories Agency (Weybridge) (Addlestone, UK). These strains have been well characterized in *in vitro* studies (Walker *et al.*, 1999) and in chicken models (Cooper *et al.*, 1990) in this laboratory. Strain 27655R was kindly provided by Professor T. Wadström (Lund University, Sweden) and was originally isolated as a rough *S. enteritidis* from a human patient suffering from gastroenteritis (Baloda *et al.*, 1988). We have reported recently the 'atypical' expression of SEF17 by 27655R amongst *S. enteritidis* (Dibb-Fuller *et al.*, 1997; Allen-Vercocoe *et al.*, 1998). All cultures and mutants derived from these strains were maintained on Dorset's egg slopes at 4 °C.

Cultural conditions. For studying the expression of fimbriae and flagella and their subsequent effect in tissue culture models, bacteria were grown under cultural conditions permissive of elaboration of each appendage. Expression of SEF14, SEF17, SEF21 and flagella has previously been reported (Thorns *et al.*, 1992; Dibb-Fuller *et al.*, 1997; Sojka *et al.*, 1996; Hotani, 1971, respectively) and was induced by growth in Sensitest agar (CM409; Oxoid) for 24 h at 37 °C, colonization factor antigen (CFA) agar (Evans *et al.*, 1977) for 72 h at 18 °C, heart infusion broth (HIB) (0038-17-7; Difco) for 48 h at 37 °C, and nutrient broth (CM1; Oxoid) for 24 h at 37 °C, respectively. All cultures were incubated statically in aerobic environments, unless otherwise stated.

Production of purified reagents and specific antibodies. Purification of SEF17, SEF21 and flagella, and the production

of monoclonal and polyclonal antibodies, have been described previously (Collinson *et al.*, 1991; Sojka *et al.*, 1996; Hotani, 1971). Soluble fibronectin and LPS were commercial reagents (F-0895 and L-4774, respectively; Sigma).

Direct binding ELISA. Cultures were grown under conditions permissive of expression of surface appendages. After pelleting by centrifugation at 2500 g for 10 min, bacteria were resuspended in 0.1 M carbonate buffer pH 9.6 and the suspension adjusted to give an optical density of 1.00 at 540 nm. Suspensions were coated on Microtitre plates (Maxisorb; Nunc), dried overnight at 37 °C and washed four times in 0.1 M PBS (12.85 g Na₂HPO₄·2H₂O l⁻¹, 3.36 g NaH₂PO₄ l⁻¹, 85 g NaCl l⁻¹; pH 7.2) containing 0.05 % (v/v) Tween 20. The washing step was repeated after each incubation step and all subsequent incubations were performed at 37 °C for 1 h except for the substrate, which was performed at room temperature for 5 min. The free binding sites were blocked with 3 % (w/v) dried milk (Marvel) in PBS (200 µl). All other ELISA reagents were added at 100 µl per well. mAb was added, followed by goat anti-mouse horseradish-peroxidase-conjugated antibody (Sigma) and then tetramethylbenzidine (TMB; Cambridge Veterinary Sciences) substrate. The reaction was stopped with 10 % (v/v) sulphuric acid and absorbance recorded at 450 nm.

Construction of mutants. Construction of isogenic mutants of *S. enteritidis* S1400/94 was described previously (Allen-Vercocoe & Woodward, 1999). Single mutants in strain LA5 and 27655R or multiple mutants were produced by P22 transduction following the methods of Anderson *et al.* (1996).

Association and invasion assays. Tissue culture cells were maintained in Eagle's Modified Essential Media (EMEM; Sigma) supplemented with 10 % heat-inactivated foetal calf serum (Sigma), 2 mM L-glutamine (Sigma) and 50 µg gentamicin (Sigma) ml⁻¹.

Confluent monolayers of human intestinal epithelial cell lines INT-407 and Caco-2 (ECACC; Salisbury) were prepared in 24-well tissue culture plates (Nunc) and pre-washed twice in Hanks' balanced salt solution (HBSS; Sigma) prior to association and invasion assays. Confluent monolayers contained between 4 and 6 × 10⁵ cells per well, and three wells were used for each test. For the association assay, INT-407 cells were infected at a ratio of 1:100 with pre-washed bacteria (in EMEM) and incubated for 2 h at 37 °C (5 % CO₂). Monolayers were washed six times with HBSS before disrupting with 1 % (v/v) Triton (Sigma). C.f.u. were determined by serial dilutions plated onto Brilliant Green agar (CM329; Oxoid). Association is reported as a combination of adhering and invading bacteria.

Invasion assays were performed in duplicate 24-well plates. After allowing the bacteria to associate with the monolayer, wells were washed three times before adding EMEM containing 100 µg gentamicin ml⁻¹. Plates were incubated at 37 °C (5 % CO₂) for 2 h and washed twice. The monolayers were then disrupted with Triton and c.f.u. were determined as described above.

Each association and invasion assay was performed at least twice (three wells per test) and representative results are reported. For statistical analyses, counts were transformed to their logarithm to base 10, and analyses of variance (ANOVA) performed, followed by standard *F*-tests.

Blocking assays. Mannose, polyclonal antisera (1/100 dilution) and mAb (1 mg ml⁻¹) were added to the bacterial suspension and the purified fimbriae, flagella, LPS and fibronectin added to the tissue culture cells. All reactions were

incubated for 15 min at 37 °C (5% CO₂) with gentle agitation prior to infection of the monolayer. Counts were statistically compared by one-way ANOVA.

Histological staining of infected INT-407 cells. Confluent monolayers prepared in six-well dishes (Bibby) on 18 mm diameter round coverslips were washed twice in HBSS and infected with 2.5×10^8 bacteria in EMEM (5 ml). After incubating for 2 h at 37 °C (5% CO₂), monolayers were washed six times in HBSS, fixed for 10 min with cold methanol (70%) and stained with 10% Giemsa stain (Sigma) for 1 h. Coverslips were washed four times in deionized water and differentiated in 1% acetic acid (1 min), washed again, air-dried and mounted onto slides with DPX mountant (Sigma). Slides were examined by light microscopy under oil immersion ($\times 1000$ magnification). Adherent cells were visualized with a Zeiss light microscope (Axiovert 25) equipped with a Contax 167MT camera.

RESULTS

Production of single and multiple mutants

Single defined mutants were prepared in *S. enteritidis* strain S1400/94 by insertion of antibiotic resistance cassettes within target genes as previously described (Allen-Vercocoe & Woodward, 1999). Construction of single and multiple mutants in strains LA5 and 27655R was made as required by P22-mediated transduction with selection for the internal marker (Table 1). All transductants were quantitatively assayed for the expression of SEF14, SEF17, SEF21 and flagella by direct binding ELISA (Table 1). Mutants failed to elaborate the

appendage encoded by the insertionally inactivated gene. No evidence was gained that mutation of any one locus altered expression of any other surface appendage.

Fimbriae- and flagella-mediated association with and invasion of INT-407 monolayers

Wild-type and single mutants were assessed for association and invasion in the INT-407 model. In all three strains with the SEF14 mutation there was no significant reduction in association or invasion of tissue culture cells. For all strains, the wild-type associated in greater numbers than the isogenic SEF17 mutants ($P < 0.001$), whilst for bacterial invasion the differences were significant for LA5 ($P = 0.02$) and S1400/94 ($P = 0.659$), and highly significant for 27655R ($P < 0.001$). Wild-type associated ($P < 0.001$) and invaded ($P < 0.001$) in significantly higher numbers than the SEF21 mutants in all three strains. Also, aflagellate mutants associated and invaded significantly less ($P < 0.001$) than their respective wild-types. In all experiments where it was tested, strain 27655R associated in significantly greater numbers than LA5 and S1400/94 (Fig. 1a, b, c).

Multiple mutants prepared in LA5 were tested in the INT-407 and Caco-2 models (Fig. 2). Counts following infection of INT-407 with LA5 wild-type were compared statistically with single and multiple mutants using one-way ANOVA. Association of the wild-type was significantly higher than all the mutants except for B214 (SEF14⁻), with the significance probability for EAV12

Table 1. Direct binding ELISA using specific mAbs to determine expression of fimbriae and flagella by single and multiple mutants of *S. enteritidis* when grown under permissive conditions

Strain	Reference name	Genotype	Antibiotic marker*	Phenotype†			
				SEF14	SEF17	SEF21	Flagella
LA5	Wild-type			+++	+	+	++
	B214	<i>sefA</i>	Kan	—	+	+	++
	EAV12	<i>agfA</i>	Bla	+++	—	+	++
	EAV3	<i>fimD</i>	Tet	+++	+	—	++
	EAV10	<i>fliC</i>	Cam	+++	+	+	—
	EAV42	<i>sefA agfA</i>	Kan Bla	—	—	+	++
	EAV21	<i>sefA agfA fimD</i>	Kan Bla Tet	—	—	—	++
	EAV40	<i>sefA agfA fimD fliC</i>	Kan Bla Tet Cam	—	—	—	—
S1400/94	Wild-type			+++	+	+	++
	EAV13	<i>sefA</i>	Kan	—	+	+	++
	EAV11	<i>agfA</i>	Bla	+++	—	+	++
	EAV1	<i>fimD</i>	Tet	+++	+	—	++
	EAV9	<i>fliC</i>	Cam	+++	+	+	—
				+++	+	+	—
27655R	Wild-type			+++	++	+	++
	EAV28	<i>sefA</i>	Kan	—	++	+	++
	EAV30	<i>agfA</i>	Bla	+++	—	+	++
	EAV29	<i>fimD</i>	Tet	+++	++	—	++
				+++	++	—	++

* Kan, kanamycin; Bla, ampicillin; Tet, tetracycline; Cam, chloramphenicol.

† A₄₅₀ in direct binding ELISA: + + +, > 1.0; + +, > 0.5; +, > 0.1; —, < 0.1.

Fig. 1. Association (□) and invasion (■) of INT-407 monolayers by *S. enteritidis* wild-type and single mutants. Bacteria were cultured under conditions permissive for expression of appendages prior to infection of monolayers. Results are expressed as c.f.u. per well $\pm$ SE. (a) SEF14, (b) SEF17, (c) SEF21, (d) flagella.

Fig. 2. Association (□) and invasion (■) of INT-407 (a) and Caco-2 (b) monolayers by single and multiple mutants of *S. enteritidis*. Bacteria were cultured in HIB for 48 h prior to infection of monolayers. Results are expressed as c.f.u. per well $\pm$ SE. LA5, wild-type; B214, SEF14⁻; EAV12, SEF17⁻; EAV3, SEF21⁻; EAV10, Fla⁻; EAV21, SEF14⁻ SEF17⁻ SEF21⁻; EAV40, SEF14⁻ SEF17⁻ SEF21⁻ Fla⁻.

(SEF17⁻) being $P = 0.025$ and the remaining differences being $P < 0.001$. EAV10 (Fla⁻) and EAV40 (SEF14⁻ SEF17⁻ SEF21⁻ Fla⁻) exhibited significant reductions in invasion ($P < 0.001$) compared to wild-type.

The trend of association and invasion of single and multiple mutants in Caco-2 cells was very similar to INT-407 (Fig. 2).

Blocking of fimbriae- and flagella-mediated association and invasion

Purified antigens and specific antibodies were incorporated into the INT-407 model to specifically block the interaction between bacteria and epithelial cells (Table 2a, b, c). The addition of polyclonal antisera and purified SEF17 fimbriae significantly reduced association ($P <$

Table 2. Blocking of fimbriae- and flagella-mediated association and invasion of INT-407 monolayers using neutralizing antibodies and purified surface antigens

Results are expressed as c.f.u. per well $\pm$ SE. **, Significant difference from wild-type ($P = 0.021$); **, significant difference from wild-type ($P < 0.001$); no asterisks, no significant difference from wild-type.

Strain†	Blocking agent	Association	Invasion		
(a) SEF17-mediated association and invasion					
27655R	None	8.75 ± 0.38 × 10 ⁷	1.55 ± 0.09 × 10 ⁶		
	SEF17 polyclonal antisera (chicken)	1.75 ± 0.20 × 10 ^{7**}	9.00 ± 1.79 × 10 ^{5**}		
	Soluble fibronectin (100 µg)	7.50 ± 1.30 × 10 ⁷	1.45 ± 0.20 × 10 ⁶		
	Purified SEF17 fimbriae (100 µg)	1.65 ± 1.20 × 10 ^{7**}	1.30 ± 0.16 × 10 ⁶		
	<i>S. enteritidis</i> LPS (100 µg)	7.25 ± 1.00 × 10 ^{7*}	1.55 ± 0.15 × 10 ⁶		
EAV30 (SEF17 ⁻)	None	1.20 ± 0.16 × 10 ^{7**}	6.50 ± 1.20 × 10 ^{4**}		
Strain†	Blocking agent	Association	Invasion		
(b) SEF21-mediated association and invasion					
LA5	None	1.60 ± 0.14 × 10 ⁶	1.90 ± 0.42 × 10 ⁵		
	D-Mannose (1 %)	2.05 ± 0.21 × 10 ^{5**}	1.30 ± 0.24 × 10 ⁵		
	mAb M8614 (1 mg)	1.70 ± 0.11 × 10 ⁶	9.90 ± 1.30 × 10 ⁴		
	mAb M8642 (1 mg)	1.90 ± 0.15 × 10 ⁶	8.10 ± 3.50 × 10 ⁴		
	mAb M8732 (1 mg)	2.05 ± 0.23 × 10 ^{5**}	9.90 ± 1.50 × 10 ⁴		
	mAb M8743 (1 mg)	1.85 ± 0.13 × 10 ⁶	1.00 ± 0.10 × 10 ⁵		
	SEF21 polyclonal antisera (chicken)	2.10 ± 0.20 × 10 ⁶	1.30 ± 0.20 × 10 ⁵		
	Purified SEF21 fimbriae (100 µg)	1.70 ± 0.19 × 10 ⁶	1.22 ± 0.14 × 10 ⁵		
	<i>S. enteritidis</i> LPS (100 µg)	1.65 ± 0.14 × 10 ⁶	1.15 ± 0.21 × 10 ⁵		
EAV3 (SEF21 ⁻)	None	2.15 ± 0.01 × 10 ^{5**}	1.30 ± 0.20 × 10 ⁵		
Strain†	Blocking agent	Without centrifugation		With centrifugation‡	
		Association	Invasion	Association	Invasion
(c) Flagella-mediated association and invasion					
LA5	None	1.10 ± 0.18 × 10 ⁶	1.70 ± 0.15 × 10 ⁴	1.14 ± 0.11 × 10 ⁶	2.00 ± 0.14 × 10 ⁴
	Purified flagella (1 mg)	1.30 ± 0.16 × 10 ⁶	2.20 ± 0.23 × 10 ⁴	1.60 ± 0.09 × 10 ⁶	2.40 ± 0.32 × 10 ⁴
	Flagella polyclonal sera	1.50 ± 0.05 × 10 ⁶	2.40 ± 0.17 × 10 ⁴	1.95 ± 0.84 × 10 ⁶	2.30 ± 0.22 × 10 ⁴
EAV19 (Fla ⁻)	None	1.30 ± 0.19 × 10 ^{5**}	1.70 ± 0.13 × 10 ^{3**}	1.60 ± 0.14 × 10 ^{5**}	1.40 ± 0.18 × 10 ^{4**}

† Bacteria were cultured in CFA agar for 24 h at 30 °C (a); in HIB for 48 h at 37 °C (5% CO₂) (b); and in nutrient broth for 24 h at 37 °C (c).

‡ Bacteria centrifuged onto monolayer at 250 g for 10 min.

0.001) as did LPS, less significantly ($P = 0.021$), but association was unaffected by the addition of soluble fibronectin. Polyclonal sera significantly reduced bacterial invasion ($P < 0.001$), whereas the other treatments had no effect.

SEF21-mediated association was significantly reduced by the addition of D-mannose and mAb M8732 ($P < 0.001$), whereas invasion was unaffected by any of the reagents added.

Centrifugation of wild-type *S. enteritidis* onto the monolayer increased bacterial association by a small, but highly significant amount ($P < 0.001$) and by two-

fold with EAV10 (aflagellate mutant). Also, a significant increase in invasion ($P < 0.001$) was observed after centrifugation of the aflagellate mutant. The addition of purified flagella or polyclonal sera had no effect.

Histological examination of infected monolayers

Wild-type 27655R adherence was localized and often aggregative (Fig. 3a), whereas wild-type LA5 cultured to express SEF21 colonized the INT-407 cells in a random, diffuse pattern (Fig. 3c). EAV12 (27655R, SEF17⁻) and EAV3 (LA5, SEF21⁻) associated visibly less than wild-types (Fig. 3b and d, respectively).

Fig. 3. Adhesion of *S. enteritidis* to INT-407 cells. 27655R (a) and EAV12 (SEF17⁻) (b) were cultured on CFA agar for 24 h at 30 °C, and LA5 (c) and EAV3 (SEF21⁻) (d) were cultured at 37 °C in HIB for 48 h (5% CO₂) before infecting monolayers. Bars, 10 µm.

DISCUSSION

To elucidate the role of fimbriae and flagella in colonization and invasion of transformed human intestinal epithelial cells, a panel of single and multiple mutants was prepared in three strains of *S. enteritidis* by insertional inactivation of target genes. The loss of phenotypic expression was confirmed by direct binding ELISA after culture of mutants under conditions permissive for elaboration of appendages. Expression of the surface appendages encoded by undisrupted genes remained unaffected.

Strain 27655R consistently associated with INT-407 cells significantly more than LA5 and S1400/94 (Fig. 1), which confirmed the findings of Baloda *et al.* (1988). We have recently reported the 'atypical' expression of SEF17 by 27655R amongst *S. enteritidis* (Dibb-Fuller *et al.*, 1997), although aberrant elaboration of this fimbrial structure was not attributed to a frameshift mutation in the *rpoS* global regulatory gene harboured by this strain (Allen-Vercoe *et al.*, 1998). The structural fimbrial subunit of SEF17, AgfA, is genetically and antigenically related to *E. coli* curli fimbriae (Collinson *et al.*, 1992) and thin aggregative fimbriae of *S. typhimurium* (Stolpe *et al.*, 1994). Expression of thin aggregative fimbriae of *S. typhimurium* has been demonstrated to promote interaction with mouse small intestinal epithelial cells (Sukupolvi *et al.*, 1997). These results are consistent with the behaviour of 27655R in our model. Association of the *agfA* mutant was significantly reduced and histological examination suggested a role for SEF17 in localized, often aggregative adherence, which could be specifically blocked with polyclonal sera or purified fimbriae (Table 2a). SEF17 would appear to promote an extremely aggregative phenotype, but it is unclear whether SEF17 mediated aggregation in the inoculum, although macroscopic clumping in cultures was not observed. An additional role in biofilm formation has been suggested by Austin *et al.* (1998), which supports the concept that SEF17 mediates aggregation.

Type 1 fimbriae of *S. typhimurium* and some *E. coli* are regarded as virulence determinants which play a role in colonization of gut epithelia (Lockman & Curtiss 1992; Takeuchi, 1967; Vidotto *et al.*, 1997) and invasion (Ernst *et al.*, 1990). However, Elliot & Kaper (1997) reported that type 1 fimbriae had no effect on levels or patterns of adhesion to cultured human cells in enteropathogenic *E. coli* (EPEC) infections. Our results suggest SEF21 mediated association in all three strains, although to observe this it was necessary to induce expression of fimbriae prior to infection (Fig. 1). Thus, a correlation was observed between level of association and SEF21 expression in the inoculum as measured by MSHA. These results agree with those of Tavendale *et al.* (1983).

SEF14 mutants in all three strains reacted very similarly, with no significant difference in either association or invasion compared to their respective wild-type. These results are consistent with those of Thorns *et al.* (1996) and Oginni *et al.* (1997). Interestingly, a role for SEF14

in transovarian transmission in chickens has been suggested by Thiagarajan *et al.* (1996).

Aflagellate mutants of *S. enteritidis* associated and invaded significantly less than motile bacteria, and numbers of associated bacteria were increased only marginally by centrifugation of bacteria onto monolayers. Numbers of associating bacteria were unaffected by the addition of purified flagella or specific polyclonal antibody to the flagellated wild-type. It is highly likely that flagella assist colonization of epithelial cells by enabling motility rather than providing an adhesin. This confirms our earlier work with adherence to chick gut explants (Allen-Vercoe *et al.*, unpublished results). Interestingly, Jones *et al.* (1992) reported that the direction of flagellar rotation affected the ability of *S. typhimurium* to invade cultured epithelial cells and, in addition, flagella have been implicated in assisting survival of *S. typhimurium* within murine macrophages (Weinstein *et al.*, 1984).

Characterization of fimbriae- and flagella-mediated interaction with cultured epithelial cells is an initial step in evaluating the role of these surface structures in pathogenesis of *S. enteritidis*. Tissue culture models have limitations for they cannot recreate the gut environment but this study demonstrates the potential role of SEF21 and SEF17 in adherence, and suggests a motility function for flagella in an undoubtedly multifactorial process. The question arises as to whether these observations will prove consistent in studies with chick gut explant models and chick *in vivo* studies.

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