MICROBIAL PATHOGENICITY

Preferential adherence of cable-piliated 
*Burkholderia cepacia* to respiratory epithelia of 
CF knockout mice and human cystic fibrosis lung 
explants

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The *Burkholderia cepacia* complex consists of at least five well-documented bacterial 
 genomovars, each of which has been isolated from the sputum of different patients with 
 cystic fibrosis (CF). Although the world-wide prevalence of this opportunistic pathogen in 
 CF patients is low (1–3%), ‘epidemic’ clusters occur in geographically isolated regions. 
 Prevalence in some of these clusters is as high as 30–40%. The majority of CF *B. cepacia* 
 isolates belong to genomovar III, but the relationship between genomovar and 
 virulence has not yet been defined. Because the initial stage of infection involves 
 bacterial binding to host tissues, the present study investigated differences in the binding 
 of representative isolates of all five genomovars to fixed nasal sections of UNC *cfr* (−/−) 
 and (+/+) mice and to human lung explants, biopsy and autopsy tissue of CF and non-CF 
 patients. Binding was highest for isolates of genomovar III, subgroup RAPD type 2, 
 but only if the isolates expressed the cable pili phenotype. Antibodies to the 22-kDa 
 adhesin of cable pili virtually abolished binding. Binding occurred only to *cfr* (−/−) 
 nasal sections or to CF lung sections and was negligible in *cfr* (+/+) or human non-CF, 
 histologically normal lung sections. Unlike normal epithelia, the hyperplastic epithelia of 
 CF bronchioles were enriched in cytokeratin 13, a 55-kDa protein that has previously 
 been shown to act as a receptor *in vitro* for cable-piliated *B. cepacia*. These findings may 
 help to explain the high transmissibility of Cbl-positive, genomovar III strains of *B. 
 cepacia* among CF patients.

Introduction

*Burkholderia cepacia* was described originally as a 
 phytopathogen [1], but has been recognised since the 
 1970s as an opportunistic respiratory pathogen in cystic 
 fibrosis (CF) patients [2–5]. It is also seen in some 
 patients with immunosuppression and chronic granulo-
 matous disease [6, 7]. In CF patients infection with *B. 
 cepacia* tends to occur in ‘epidemic’ geographic 
 clusters, with regional differences in prevalence. In 
 Canada as a whole, the prevalence is 8–10% [8], but in 
 the Toronto adult CF clinic prevalence is much higher 
 (46%) [9]. Patients who become colonised show 
 unpredictable and variable clinical responses, ranging 
 from no apparent deleterious consequences to a rapidly 
 fatal necrotising pneumonia and septicaemia (the 
 cepacia syndrome). The same variability of outcome 
 has been noted in other centres, suggesting that only 
 some CF patients are susceptible to serious respiratory 
 disease with this pathogen. However, large national 
 surveys indicate that patients infected with *B. cepacia* 
 exhibit an overall adverse clinical course, with higher 
 mortality rates than those colonised only with *Pseu-
 domonas aeruginosa*, the more common opportunist 
 pathogen in CF [4, 10–13]. In particular, *B. cepacia*- 
 infected CF patients have a much higher post-operative 
 mortality rate (64% versus 19%) in the first 3 months 
 following double lung transplantation [14]. Because the 
 organism is generally resistant to antibiotics and other 
 forms of therapy, there is a pressing need to discover 
 more about the mechanisms of its pathogenesis, 
 including the colonisation process, virulence properties 
 and mechanisms of drug resistance.

*B. cepacia* isolates from sputum of CF patients and 
 from environmental sources (plants, soil) are geneti-
cally diverse and have been categorised into at least five genomic groups (genomovars), based on computer-assisted cluster analyses of whole-cell protein profiles, fatty acid content and DNA–DNA and DNA–rRNA hybridisation assays [15]. The major ‘epidemic’ clusters affecting Canadian CF patients are associated with genomovar III. Less frequently, genomovars II (B. multivorans) and IV have been found. The rapid spread of genomovar III from patient to patient attests to its high index of transmissibility. Environmental strains tend to belong to genomovar I and are rarely cultured from CF patients.

The relationship between genomovar and virulence has not yet been defined. Potential virulence determinants of B. cepacia include a potent haemolysin [16], a siderofer [17], LPS [18, 19], an extracellular protease [20], lipases [21], a melanin-like pigment [22] and surface cable pili. The latter have been shown to mediate bacterial adherence to mucin and epithelial cells [23, 24]. It is not known which (if any) of these factors has the most negative impact on CF patients. However, because acquisition of B. cepacia tends to increase with age and duration of lung disease, virulence probably also depends in large measure on compromised host defences. The chronic stasis of mucus in the airways, impaired mucociliary clearance, altered electrolyte composition, defective CFTR, inactive defensins, epithelial metaplasia, neutrophil products and other consequences of pre-existing infection and inflammation, may all play a role in B. cepacia persistence and host susceptibility to further lung damage.

One highly transmissible clone (also known as the ET strain) is prevalent in Canadian and UK CF centres, belongs to genomovar III, RAPD type 2, and carries the ‘epidemic’ DNA marker (BCESM) [25]. This strain also carries the cbcl gene encoding the major cable pilin subunit [26, 27]. In previous studies involving a group of 15 CF isolates from Ontario, all the isolates were ET strains and 14 of them expressed the unique cable pilin phenotype [26, 28]. The pili were found to mediate specific binding to mucin carbohydrates as well as to a 55-kDa protein present in buccal epithelial cells [23, 24, 29]. Recently, the 55-kDa cell receptor has been identified as cytokeratin 13 [30]. The actual adherence mechanism involves a 22-kDa adhesin protein associated with the cable pili [23, 24].

The present study involves a test of B. cepacia adherence to fixed tissue sections of CF and control respiratory epithelia; nasal epithelia of UNC CF knockout mice [31] and human lung explant, biopsy or autopsy tissues were used. Despite the lack of overt lung pathology in UNC cftr knockout (−/−) mice, their nasal epithelia exhibit mild inflammatory lesions and chloride flux pathology that closely mimic human CF respiratory epithelia [32, 33]. The binding properties of representative isolates of B. cepacia from all five genomovars to the mouse and human tissue sections were compared, to decide whether specific genomovar strains might be considered to have a greater potential for adherence and colonisation in vivo than others. The sections were also used to ascertain if CF tissue was more susceptible than non-CF tissue to B. cepacia binding.

**Materials and methods**

**B. cepacia strains and culture conditions**

Sputum isolates BC5, BC7, BC24 and BC45 were from the B. cepacia collection held in this laboratory and have been described previously [24, 26, 28]. Isolate BC7, which was used extensively in the present study, was isolated originally from a 15-year-old CF patient who was clinically healthy until acquisition of B. cepacia. The patient died with the cepacia syndrome 1 month later. Isolate BC45 was cultured from a CF patient who was chronically infected with P. aeruginosa for 20 years, then acquired B. cepacia, but did not die until 4 years later at the age of 43. This patient never developed the cepacia syndrome. Isolate BC24 was isolated from a patient who had severe lung disease (and harboured P. aeruginosa) for several years before acquiring B. cepacia at the age of 21. He died 2 years later but without signs of the cepacia syndrome. Isolate BC5 was isolated from a 16-year-old patient with only mild symptoms of lung disease. Three years later the patient was alive, but moved out of town and was lost to follow-up. All other isolates in Table 1 were from the B. cepacia research panel of strains held by Dr E. Mahenthiralingam (Department of Pediatrics, University of British Columbia). Isolate J2315, donated by Dr J. Govan (Edinburgh), was cultured during the 1980s from a CF patient who is still alive. Species identity of all isolates was reconfirmed in this laboratory by growth on B. cepacia selective isolation agar [34]. Isolates were stored at −80°C and routinely subcultured on Brain Heart Infusion Agar (Becton Dickinson, Cockeysville, MD, USA) at 37°C for 48 h before use. Single colonies were inoculated into 10 ml of Tryptic Soy Broth (TSB; Difco) and grown at 37°C for 24 h on an orbital shaker at 150 rpm in preparation for experiments.

**Fluorescein isothiocyanate (FITC)-labelling of B. cepacia**

All B. cepacia isolates were labelled with FITC following the method of Falk et al. [35]. Labelled bacteria were recovered by centrifugation, washed three times with phosphate buffered saline (PBS) containing bovine serum albumen (BSA) 0.1% and resuspended in PBS containing BSA 1% to give 10⁶ cfu/ml. Samples were examined by light and fluorescence microscopy (Leica Dialux 22, Willowdale, Ont, Canada) to confirm that all bacteria from each isolate were fluorescent.
Either 100- or 200-μl volumes of the bacterial suspension were used in binding experiments.

Nasal epithelial sections of mice

Animal experiments were performed under protocols approved by the animal ethics committee at the Hospital for Sick Children, Toronto. Wild-type C57Bl/6J mice (Jackson Laboratory, Bar Harbor, Maine, USA) fed a normal chow diet were used as controls. UNC efr (−/−) and efr (+/+ ) mice [31] from the same colony were housed in clean conventional rooms free of rodent viral, mycoplasma, and bacterial pathogens and maintained on a liquid diet as described earlier [33]. Mice were killed between 6 and 8 weeks of age by intraperitoneal chloral hydrate (3.6%) and exsanguination. Transverse nasal sections were made 2.8 mm from the tip of the nose, nasal specimens were fixed in buffered formalin 10%, decalcified in formic acid 8% [36] and embedded in paraffin. Nasal sections (5 μ thick) were cut for haematoxylin and eosin (H and E) staining or bacterial binding assays.

In some experiments, 4 μl of a 1% solution of the amphotropic detergent polyoxyethylene-9-lauryl ether (Polidocanol, Sigma) were instilled into the right nostril of wild-type C57Bl/6J mice to induce a mild inflammatory reaction [37]. After 30 min mice were killed and nasal sections were prepared as described above. Binding of bacterial isolates was compared in the normal and inflamed nasal cavities by fluorescence microscopy.

Lung sections from CF and non-CF patients

Formalin-fixed, paraffin-embedded lung explants, biopsy or autopsy tissue were obtained from the pathology tissue library at the Hospital for Sick Children, Toronto. For each patient sample one (or more sections) were available from proximal regions (to show many large bronchioles and parenchyma) and peripheral regions (containing mainly small bronchioles and alveoli). Sections included explants of six CF patients, a biopsy section from one non-CF patient and autopsy lung sections from two non-CF patients. The lung sections of the non-CF patients showed no microscopic pathology and were used as normal controls.

Bacterial binding to tissue sections

Human lung or mouse nasal sections were deparaffi-
nised in xylene, rehydrated in graded alcohol, washed twice in PBS and blocked with BSA 3% for at least 2 h at room temperature. FITC-labelled B. cepacia (0.1 or 0.2 ml of 10^6 cfu/ml) in PBS-BSA 1% were then added and incubated for 1 h at 37°C in a humidified chamber. The slides were washed six times with PBS-BSA 0.1% to remove non-adherent bacteria, fixed in buffered formalin 10% for 15 min, washed three times with PBS, counter-stained with Mayer’s haematoxylin and mounted. Sections were examined in a fluorescence microscope and bacterial binding was quantified by counting the number of bacteria associated with mouse nasal tissue (10 fields per section) or human respiratory epithelia, as detailed in the Results section. Binding of ≈10 bacteria was considered to be non-specific background binding.

Antibody to the 22-kDa adhesin of cable pili

Immunoreactivity of isolates with a polyclonal anti-
body to the 22-kDa adhesin of cable pili was tested by immunoblots of outer-membrane protein preparations, by protocols described previously [24]. This antibody has been shown to react with a doublet band centred at 22 kDa in isolates expressing surface cable pili. To test for the involvement of cable pili in the binding of B. cepacia to lung sections, FITC-labelled isolate BC7 was pre-incubated with the antibody to the 22-kDa adhesin for 1 h at 37°C, centrifuged, resuspended in PBS-BSA 1% and incubated with lung sections as in a standard binding assay.

Detection of the cblA gene and Bcesm DNA marker in isolates

PCR amplification was used to detect the cblA gene by an approach similar to that described by Yoshi et al. [38]. Briefly, intact bacterial cells (1 × 10^6 cfu) of each isolate were used as a source of template DNA. The reaction mixture was heated at 95°C for 12 min to release DNA from the bacteria and then subjected to 30 cycles of amplification (1 min at 94°C, 1 min at 60°C, 1 min at 72°C). A typical reaction mixture consisted of the cblA gene specific primers GTCCAGAAGGACAT TACCGTC (sense primer), GGGCTTAGCTACCC GAAGCCGGGCT (antisense primer) (0.4 μM each), dNTPs (200 μM each), B. cepacia (1 × 10^6 cfu), AmpliTaq Gold DNA polymerase (2.0 U, Perkin-Elmer Cetus, Norwalk, CT, USA), MgCl2 (1.5 mM) and Tris-
HCl buffer (10 mM containing 50 mM KCl). Amplicons were analysed by electrophoresis on agarose 1.5% gel and stained with ethidium bromide to detect the expected 447-bp cblA gene product [26]. The presence of the Bcesm DNA marker in isolates was confirmed by PCR by using methods described earlier by Mahenthiralingam et al. [39].

Buccal epithelial cell fractions

Buccal epithelial cells (BEC) were fractionated to produce a cytokeratin-rich fraction by the method of Franke et al. [40]. This fraction contains the major cytokeratin (55 kDa) that acts as a receptor for Clbl-induced strains of B. cepacia in binding assays [30], and was used in the present study either as a receptor in bacterial overlay assays or as an inhibitor of bacterial binding to tissue sections. Binding of B.
cepacia isolates to blots of BEC extracts was tested by overlay of 35S-labelled bacteria as described earlier [23, 30]. For inhibition studies, isolates were pre-incubated with the BEC extract for 1 h at 37°C, centrifuged, resuspended in PBS-BSA 1% and incubated with lung sections as in a standard binding assay.

**Immunodetection of cytokeratin 13 in CF and non-CF lung sections**

Cytokeratin 13 was detected by indirect immunofluorescence with a monoclonal antibody (MAB) to cytokeratin 13 (Novocasta Laboratories) (1:100 v/v) according to the manufacturer’s instructions. Detection was by an anti-mouse IgG-Cy3 fluorophore (1:250 v/v). Sections were counterstained by Meyer’s haematoxylin.

**Results**

**Nasal histology of wild-type, cfr (−/−) and (+/+ ) mice**

H and E nasal sections taken from wild-type (chow-fed C57BL/6J) and UNC cfr (+/+ ) mice (six per group) exhibited normal morphology of olfactory, stratified squamous, sepal and turbinate epithelia. In particular, the respiratory epithelia of the septa and turbinates showed a normal distribution of pseudostratified ciliated columnar cells, goblet cells and basal cells [41]. However, in each of nine cfr (−/−) mice examined there was evidence of mild inflammation with mucosal swelling, luminal exudates, epithelial hyperplasia, patchy losses of cilia and sparse leucocyte infiltration in the lamina propria (not shown). These pathological changes were very similar to those described earlier for UNC cfr (−/−) mice [32]. In two wild-type mice given intranasal Polisoracol detergent 30 min before killing, there was evidence of mild mucosal swelling and partial denudation of surface respiratory epithelia on the treated side of the nostril.

**Binding of FITC-labelled B. cepacia to nasal epithelia**

Each isolate shown in Table 1 was labelled with FITC and separately overlaid in triplicate on fixed nasal sections prepared from four wild-type, one cfr (+/+) and five cfr (−/−) mice. Because cfr (+/+ ) mice were in short supply, and preliminary experiments revealed no differences in binding between wild-type and cfr (+/+ ) sections, wild-type nasal sections were used in most cases instead of cfr (−/−) sections. None of the isolates bound to cfr (−/−) or wild-type nasal sections, but binding of a subset of the genovar III, RAPD type 2 isolates (BC5, BC7, BC24, C5424) was prominent on the epithelium and luminal exudates of cfr (−/−) sections, as shown for one of the isolates in Fig. 1. Table 2 shows representative results for isolate BC7 after counting the number of bacteria bound to the epithelium in 10 randomised fields (0.4 × 0.4 mm) per section. Isolates BC5, BC24 and C5424 gave similar results. Isolates BC45 and J2315, which also belong to genovar III, RAPD type 2, did not bind and were indistinguishable from the environmental isolate ATCC 25416 (<10 bacteria per field) and all other isolates.

As the mildly inflamed respiratory epithelia of cfr (−/−) mice appeared to be much more receptive to bacterial binding than wild-type or cfr (−/−) nasal epithelium, an attempt was made to mimic these conditions by inducing a mild inflammation in the nares of wild-type mice by brief pre-treatment with detergent. However, as shown in Table 2, binding by isolate BC7 was not enhanced, indicating that an acute inflammatory reaction was insufficient to render the tissues suitable for bacterial binding. Presumably the more chronic inflammatory condition found in cfr

<table>
<thead>
<tr>
<th>Isolate no.</th>
<th>Geographic location</th>
<th>Source</th>
<th>Genomovar</th>
<th>RAPD</th>
<th>BCESM</th>
<th>cblA gene</th>
<th>22 kDa adhesin</th>
</tr>
</thead>
<tbody>
<tr>
<td>ATCC 25416</td>
<td>ATCC</td>
<td>Onion</td>
<td>I</td>
<td>039</td>
<td>–</td>
<td>–</td>
<td>–</td>
</tr>
<tr>
<td>CEP 509</td>
<td>Australia</td>
<td>CF</td>
<td>I</td>
<td>041</td>
<td>–</td>
<td>–</td>
<td>–</td>
</tr>
<tr>
<td>C5391</td>
<td>Toronto</td>
<td>CF</td>
<td>II</td>
<td>003-BM</td>
<td>–</td>
<td>–</td>
<td>–</td>
</tr>
<tr>
<td>JTC</td>
<td>Vancouver</td>
<td>CGD</td>
<td>II</td>
<td>BM-X</td>
<td>–</td>
<td>–</td>
<td>–</td>
</tr>
<tr>
<td>BC5</td>
<td>Toronto</td>
<td>CF</td>
<td>III</td>
<td>002</td>
<td>+</td>
<td>–</td>
<td>–</td>
</tr>
<tr>
<td>BC7</td>
<td>Toronto</td>
<td>CF</td>
<td>III</td>
<td>002</td>
<td>+</td>
<td>–</td>
<td>–</td>
</tr>
<tr>
<td>BC24</td>
<td>Toronto</td>
<td>CF</td>
<td>III</td>
<td>002</td>
<td>+</td>
<td>+</td>
<td>–</td>
</tr>
<tr>
<td>BC45</td>
<td>Toronto</td>
<td>CF</td>
<td>III</td>
<td>002</td>
<td>+</td>
<td>–</td>
<td>–</td>
</tr>
<tr>
<td>C5424</td>
<td>Vancouver</td>
<td>CF</td>
<td>III</td>
<td>002</td>
<td>+</td>
<td>+</td>
<td>–</td>
</tr>
<tr>
<td>J2315</td>
<td>UK</td>
<td>CF</td>
<td>III</td>
<td>002</td>
<td>+</td>
<td>–</td>
<td>–</td>
</tr>
<tr>
<td>C4415</td>
<td>Toronto</td>
<td>CF</td>
<td>III</td>
<td>006</td>
<td>–</td>
<td>–</td>
<td>–</td>
</tr>
<tr>
<td>C1394</td>
<td>UK</td>
<td>CF</td>
<td>III</td>
<td>013</td>
<td>–</td>
<td>–</td>
<td>–</td>
</tr>
<tr>
<td>PC184</td>
<td>OH, USA</td>
<td>CF</td>
<td>III</td>
<td>017</td>
<td>–</td>
<td>–</td>
<td>–</td>
</tr>
<tr>
<td>LG614291</td>
<td>Belgium</td>
<td>CF</td>
<td>IV</td>
<td>gIV-X</td>
<td>–</td>
<td>–</td>
<td>–</td>
</tr>
<tr>
<td>PC254</td>
<td>WA, USA</td>
<td>CF</td>
<td>V</td>
<td>008</td>
<td>–</td>
<td>–</td>
<td>–</td>
</tr>
</tbody>
</table>

Genomovar status of all isolates was determined by Vandamme et al. [15]. RAPD typing was done by Mahenthiralingam et al. [39]. The BCESM and cblA gene were detected by PCR as described in Materials and methods. The 22-kDa adhesin was detected by Western blot analysis of outer-membrane preparations with the anti-adhesin antibody.
Fig. 1. Binding of isolates BC7 and ATCC 25416 to nasal sections of cfr (−/−) and wild-type mice. Nasal sections from cfr (−/−) and wild-type mice were incubated with FITC-labelled isolate BC7 or ATCC 25416 (0.1 ml of 1 × 10^6 cfu/ml) for 1 h at 37°C. Sections were washed to remove unbound bacteria, fixed in buffered formalin 10% and counter-stained with haematoxylin. Airway epithelia and lumens are represented by ‘E’ and ‘L’, respectively. (a) and (b) isolate BC7 binding to cfr (−/−) and wild-type mouse nasal sections respectively; (c) ATCC 25416 binding to cfr (−/−) mouse nasal section.

Table 2. B. cepacia (isolate BC7) binding to mouse nasal sections

<table>
<thead>
<tr>
<th>Mouse type</th>
<th>Mean (SEM) number of bacteria bound</th>
</tr>
</thead>
<tbody>
<tr>
<td>cfr (+/+ )</td>
<td>2.6 (2.0)</td>
</tr>
<tr>
<td>Wild-type</td>
<td>2.7 (2.0)</td>
</tr>
<tr>
<td>Wild-type</td>
<td>4.0 (2.4)</td>
</tr>
<tr>
<td>Wild-type</td>
<td>3.7 (1.7)</td>
</tr>
<tr>
<td>cfr (−/−)</td>
<td>95.9 (12.8)</td>
</tr>
<tr>
<td>cfr (−/−)</td>
<td>75.3 (8.9)</td>
</tr>
<tr>
<td>cfr (−/−)</td>
<td>82.2 (3.5)</td>
</tr>
<tr>
<td>cfr (−/−)</td>
<td>43.0 (7.1)</td>
</tr>
<tr>
<td>cfr (−/−)</td>
<td>22.0 (2.5)</td>
</tr>
<tr>
<td>Wild-type (detergent)</td>
<td>2.6 (2.0)</td>
</tr>
<tr>
<td>Wild-type (detergent)</td>
<td>2.1 (1.5)</td>
</tr>
</tbody>
</table>

FITC-labelled isolate BC7 was incubated in triplicate with fixed sections of wild-type, UNC cfr (+/+) or cfr (−/−) mice or wild-type mice pre-treated with detergent 30 min before killing. After washing to remove non-bound bacteria, bound bacteria were counted in 10 fields and the mean and SEM were calculated for each mouse section. <10 adherent bacteria per field was considered background, non-specific binding.

Table 2. B. cepacia (isolate BC7) binding to mouse nasal sections

Histology of human lung sections

Experiments similar to those described for nasal sections were undertaken with human CF lung explants and non-CF biopsy or autopsy tissue. There were six CF patients (aged 14–35) of whom the first four (Table 3) were colonised by both B. cepacia and P. aeruginosa. Patients 5 and 6 were colonised only by P. aeruginosa. Lung sections obtained from three non-CF individuals (not shown) were histologically normal and were used for comparison.

The CF lung sections (two-to-five per patient) included proximal and distal regions, and contained large and small bronchioles, as well as alveoli. Most bronchiolar airways were enlarged, filled with mucopurulent exudates and lined by hyperplastic epithelia, or epithelia undergoing squamous metaplasia. There was partial loss of cilia, patchy exfoliation of cell layers and dense peribronchial infiltrations of inflammatory cells. Terminal and respiratory bronchioles, which are normally devoid of goblet cells in healthy lungs, contained numerous goblet cells. In alveolar regions, emphysema, thickened interstitium with mild fibrosis and inflammatory cells in the air spaces were observed. The pathological changes resembled those described earlier by Baltimore et al. [42] for CF patients infected by P. aeruginosa. Occasionally regions of more normal looking thin-walled alveoli (designated ‘thin’ in Table 3) were also recognised. Morphological differences could not be distinguished between patients infected with P. aeruginosa alone versus P. aeruginosa plus B. cepacia.
Table 3. Bound bacteria in CF lung explants

<table>
<thead>
<tr>
<th>Patient no.</th>
<th>Bronchial epithelium</th>
<th>Luminal exudates</th>
<th>Alveolar septa</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>large</td>
<td>small</td>
<td>(large bronchioles)</td>
</tr>
<tr>
<td>1</td>
<td>61.5 ± 6.1</td>
<td>21.3 ± 2.3</td>
<td>136 ± 12.5</td>
</tr>
<tr>
<td>2</td>
<td>76.2 ± 5.5</td>
<td>14.5 ± 2.0</td>
<td>144 ± 11.7</td>
</tr>
<tr>
<td>3</td>
<td>61.3 ± 6.9</td>
<td>19.4 ± 2.7</td>
<td>166 ± 7.9</td>
</tr>
<tr>
<td>4</td>
<td>56.6 ± 7.9</td>
<td>25.2 ± 1.8</td>
<td>158 ± 12.0</td>
</tr>
<tr>
<td>5</td>
<td>65.6 ± 6.6</td>
<td>12.5 ± 2.1</td>
<td>146 ± 8.1</td>
</tr>
<tr>
<td>6</td>
<td>59.4 ± 7.9</td>
<td>18.8 ± 3.1</td>
<td>113 ± 11.6</td>
</tr>
<tr>
<td>Mean (SEM)</td>
<td>63.4 ± 5.6</td>
<td>18.6 ± 1.2</td>
<td>144 ± 16.2</td>
</tr>
</tbody>
</table>

FITC-labelled isolate BC7 was incubated with fixed sections of CF lung explants. Non-bound bacteria were removed by washing, and the number of bound bacteria was counted per 100-μm length of epithelium or 100 μm² of exudate in 6–18 (average 10) fields. Values represent mean and SEM. Thick versus thin alveolar septa refer to hyperplastic, swollen versus normal septa, respectively. Large bronchioles include small bronchi and regular bronchioles; small bronchioles refer to terminal and respiratory bronchioles. All six patients were colonised by P. aeruginosa, and patients 1–4 were also colonised by *P. cepacia* at the time of double lung transplant.

**Binding of *P. cepacia* isolates to human lung sections**

Binding of *P. cepacia* isolates (listed in Table 1) to CF and non-CF lung sections was quantified by counting the number of bacteria bound per 100-μm lengths of epithelium or per 100 μm² of luminal exudate in an average of 10 fields per section. As observed with the mice nasal sections, only isolates BC5, BC7, BC24 and C5424 exhibited significant binding (>10 bacteria per field), and they bound only to CF lung sections. Representative results for isolate BC7 in CF and non-CF bronchioles are shown in Fig. 2. Quantification of

Fig. 2. Binding of isolate BC7 to CF and non-CF bronchioles. Sections of CF and non-CF lungs were incubated with FITC-labelled isolate BC7 (0.2 ml of 1 × 10⁵ cfu/ml) for 1 h at 37°C. Sections were washed, fixed in buffered formalin 10% and counter-stained with haematoxylin. Airway epithelia and lumens are represented by ‘E’ and ‘L’ respectively. (a) and (c) haematoxylin-stained sections of CF and non-CF bronchioles respectively; (b) and (d) isolate BC7 binding to sections corresponding to (a) and (c).
binding to different CF lung structures of each patient is presented in Table 3. The isolates that exhibited binding bound strongly to hyperplastic, damaged CF bronchiolar epithelia (especially in the larger bronchi-oles), to purulent exudates in the lumen and to thickened alveolar septa. There was no binding to vascular endothelia or interstitial. No differences were observed in the binding of B. cepacia to tissues obtained from patients infected with P. aeruginosa versus P. aeruginosa plus B. cepacia.

Role of the 22-kDa adhesin

Immunoblots of each isolate with the polyclonal antibody to the 22-kDa adhesin revealed that the four isolates shown earlier to be capable of binding to CF tissues contained the adhesin (Table 1 and Fig. 3), although trace amounts were also detected in isolates BC45 and J2315 (Fig. 3). The presence of the cblA gene for the Cbl pili subunit was not confined to the four high binding isolates. The other two members of the genomovar III, RAPD type 2 classification (J2315 and BC45), which did not bind to CF tissues, also carry the cblA gene (Table 1).

To assess further the role of the adhesin in the binding of B. cepacia to lung sections, isolate BC7 was pre-incubated with the anti-adhesin antibody (1 in 50 dilution) or with pre-immune serum (no inhibitor control) and binding to three CF (and one non-CF) lung sections was measured. Fig. 4 shows that the anti-adhesin caused a dramatic inhibition of binding of isolate BC7 to bronchiolar epithelium and luminal exudates of CF tissue sections. The anti-adhesin antibody had no effect on non-binding isolates. These experiments indicate that not all genomovar III isolates carrying the cblA gene have the Cbl pili phenotype, but those that do utilise the 22-kDa adhesin to bind to CF epithelial cells.

Effect of the cytokeratin extract of BEC

All the isolates shown in Table 1 were metabolically labelled with 35S-methionine and tested individually in bacterial overlay assays for their binding to the 55-kDa protein of cytokeratin extracts of BEC (Fig. 5). Isolates that belonged to genomovar III, RAPD type 2 and were cblA positive and adhesion positive (CblA+/Adh+) included BC5, BC7, BC24 and C5424 (Table 1). These isolates bound to cytokeratin 13. The two isolates (BC45 and J2315) that also belonged to genomovar III, RAPD type 2, but were cblA+/Adh− did not bind to

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**Fig. 3.** Detection of the 22-kDa adhesin in B. cepacia isolates. Outer membrane fractions from each B. cepacia isolate were electrophoresed on SDS-polyacrylamide gels (16%), transferred to nitrocellulose membranes and blocked with BSA 3%. Membranes were incubated with the anti-22-kDa adhesin antibody (1 in 1000 v/v), washed, and the bound antibody was detected by anti-rabbit IgG conjugated to alkaline phosphatase and NBT-BCIP substrate. Molecular mass standards (kDa) are shown on the left. The adhesin is a doublet centred at 22 kDa.

**Fig. 4.** Inhibition of strain BC7 binding to CF lung sections by the anti-22-kDa adhesin antibody. FITC-labelled BC7 strain cells were pre-incubated with pre-immune serum or the anti-adhesin antibody (1 in 50 dilution), centrifuged, suspended in PBS-BSA 1% and incubated with CF lung sections. Airway epithelia and lumens are represented by ‘E’ and ‘L’ respectively. (a) pre-immune serum (positive control); (b) anti-22-kDa adhesin antibody (inhibitor).
cytokeratin 13. Thus the 22-kDa adhesin plays a major role in mediating binding. No other genomovar samples bound to the 55-kDa protein or to any other component in the extracts.

Purified CK13 is not available commercially, thus its ability to inhibit binding could not be tested. However, isolate BC7 was pre-incubated with the cytokeratin-rich fraction of BEC (which contains CK13 and CK4 [43]) and then with CF lung explant sections. As shown in Fig. 6 the cytokeratin fraction caused a major reduction in binding. Bacteria that had been exposed only to the cytokeratin suspension buffer (control) were not inhibited.

Distribution of cytokeratin 13 in CF and non-CF lung sections

As only CF lung tissue exhibited susceptibility to binding by Cbl-positive *B. cepacia*, the present study tested the possibility that CF airway epithelia, especially regions exhibiting pathological hyperplasia or squamous metaplasia, might show an increased expression of cytokeratin 13. Immunofluorescent detection in CF and non-CF lungs was performed with a specific MAb that recognises cytokeratin 13. In non-CF tissues (normal bronchioles) cytokeratin 13 was not detected, as was expected from the previous distribution studies of Moll et al. [44]. However, in CF airways, cytokeratin 13 was detected in the epithelial lining of almost all small bronchi and bronchioles. CF bronchioles of relatively normal appearance contained cytokeratin 13 in the region of basal cells, whereas bronchioles exhibiting more hyperplasia and squamous metaplasia gave a fluorescent signal throughout the mucosa, i.e., in basal, middle and luminal surface regions (Fig. 7). Unfortunately, the antigen ‘unmasking’ process (heating tissue sections at 121°C for 1 min) that is part of the procedure required to detect cytokeratin 13, effectively prevented the attempts to perform colocalisation experiments of bound bacteria and cytokeratin. Nevertheless, the findings indicate that cytokeratin 13 expression is significantly increased in CF airway epithelia, and becomes apically disposed in severely affected mucosa. These are the same areas in which *B. cepacia* binding was found to be most pronounced.

Discussion

The results of this study indicate that there is a prominent functional difference among *B. cepacia*
genomovars in their ability to bind to CF airway tissue. Genomovar III strains belonging to RAPD type 2 were adherent, but within this group, only those isolates that expressed surface cable pili and the 22-kDa adhesin were able to bind. These findings extend earlier in-vitro studies which showed that cable pili mediate *B. cepacia* attachment to the isolated 55-kDa cytokeratin of BEC [23, 30]. Non-cable-pilated isolates did not bind. The two non-binding isolates (BC45 and J2315) in the RAPD type 2 group did not express the adhesin, even though they carry the *chtA* gene and are indistinguishable from the others by current DNA typing methods. Presumably a mutation in a regulatory gene or mutation(s) in one or more of the genes required for pilin assembly, transport, anchorage or polymerisation, may be responsible. However, as most of the genomovar III isolates in the research panel (and in our own published collection) do exhibit cable pili [26], the cable pili phenotype appears to be the most common genomovar III phenotype in CF patients.

Some Cbl-negative strains, particularly those belonging to genomovar II, have also been reported to be associated with ‘epidemic’ spread in CF patients living in other geographic areas [4, 12, 15]. A previous study [45] showed that genomovar III isolates with no or very few cable pili (BC45 and BC61 respectively) were able to bind to galactolipid receptors (particularly asialo GM1 and Galα3. In other genomovars were not tested in those experiments. Asialo GM1 has been reported to be increased in CF epithelial cells, and may be a candidate receptor for binding by Cbl-negative *B. cepacia*. However, as the present study failed to detect binding to CF tissues by strains belonging to other genomovars, the biological significance of in-vitro galactolipid binding remains in question. Therefore, the mechanism by which Cbl-negative strains attach to host tissue in vivo remains unsolved.

A second major finding in this study was that human CF tissues (mainly damaged epithelial linings of small airways and thickened alveolar septa) were much more susceptible to Cbl-positive *B. cepacia* binding than non-CF tissues. Even the *cfr* (−/−) mice exhibited a stronger tendency to bind the Cbl-positive strains to their nasal mucosa than *cfr* (+/+ or wild-type) mice. Apart from the absence of functional CFTPR, the only obvious feature that distinguished *cfr* (−/−) mice was the presence of an underlying mild nasal inflammation. Attempts to reproduce this state by inducing an acute mild inflammatory reaction in wild-type mice with detergent were insufficient to cause an increase in *B. cepacia* binding. Thus the chronic nature of the inflammatory process in CF, which probably entails continuous subtle damage inflicted by leukocytes or leukocyte products, may be an important prerequisite for Cbl-positive *B. cepacia* binding.

Normal epithelial tissues engaged in repair after injury undergo a remodelling of the epithelium and shifts in cytokeratin expression, with an increase in cytokeratin 13 [46]. The observed preferential attachment of Cbl-positive *B. cepacia* to cytokeratin 13 in vitro, the inhibition of binding of isolate BC7 to lung tissue by the cytokeratin fraction of BEC, and the increase in fluorescent staining of cytokeratin 13 in damaged CF airway tissues, all point to the interpretation that cytokeratin 13 plays an important receptor role for cable-pilated *B. cepacia* interactions in CF lungs.

The present studies do not prove that Cbl-positive *B. cepacia* isolates induce lung damage. However, initial binding of bacteria to host tissues is generally considered to be an important virulence determinant. As the results of the present study show that Cbl-positive isolates (cblA+/Adh+) preferentially adhere to chronically damaged CF epithelia enriched in CK13, these isolates are expected to have a greater potential than other isolates or genomovars to persist, infect and damage the CF lung. Preliminary experiments in this laboratory, with repeat nasal instillations of isolate BC7 or ATCC 25416 into *cfr* (−/−) and (+/+) mice, have
shown that after 2–3 weeks there is a much greater persistence of B. cepacia isolate BC7 in cffr(−/−) than in cffr(+/−) lungs (unpublished observations). Further research will be undertaken to extend these experiments and create an animal model suitable for studies of virulence of B. cepacia.

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