

Bifidobacterial surface-exopolysaccharide facilitates commensal-host interaction through immune modulation and pathogen protection

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Bifidobacteria comprise a significant proportion of the human gut microbiota. Several bifidobacterial strains are currently used as therapeutic interventions, claiming various health benefits by acting as probiotics. However, the precise mechanisms by which they maintain habitation within their host and consequently provide these benefits are not fully understood. Here we show that *Bifidobacterium breve* UCC2003 produces a cell surface-associated exopolysaccharide (EPS), the biosynthesis of which is directed by either half of a bidirectional gene cluster, thus leading to production of one of two possible EPSs. Alternate transcription of the two opposing halves of this cluster appears to be the result of promoter reorientation. Surface EPS provided stress tolerance and promoted *in vivo* persistence, but not initial colonization. Marked differences were observed in host immune response: strains producing surface EPS (EPS⁺) failed to elicit a strong immune response compared with EPS-deficient variants. Specifically, EPS production was shown to be linked to the evasion of adaptive B-cell responses. Furthermore, presence of EPS⁺ *B. breve* reduced colonization levels of the gut pathogen *Citrobacter rodentium*. Our data thus assigns a pivotal and beneficial role for EPS in modulating various aspects of bifidobacterial–host interaction, including the ability of commensal bacteria to remain immunologically silent and in turn provide pathogen protection. This finding enforces the probiotic concept and provides mechanistic insights into health-promoting benefits for both animal and human hosts.

The human gut is considered one of the most densely colonized ecosystems known, and is estimated to provide residence to 10–100 trillion microorganisms (1). These microbes play an important role in human nutrition and health by promoting nutrient supply, preventing pathogen colonization, and shaping and maintaining normal mucosal immunity. Indeed, these interactions work both ways, as the human gastrointestinal tract also provides the microbiota with access to key nutrients and a stable environment required for growth (2).

Bifidobacteria represent one of the dominant bacterial groups of the human intestinal microbiota (1). Certain members of the genus *Bifidobacterium* can exert specific health benefits on their host and are therefore considered to be probiotics (3). Consumption of specific bifidobacteria is associated with inhibition or reduction of cancer (4), antimicrobial activity against pathogens (5), and reduction of relapse frequency of ulcerative colitis (6). Despite these reports, molecular mechanisms underlying these health-promoting claims are largely unknown.

One of the proposed mechanisms by which bifidobacteria mediate (some of) these health benefits is the production of exopolysaccharide/capsule (EPS) (7). Bacterial EPS consists of a repeating mono- or oligosaccharide subunit connected by varying glycosidic linkages, thereby generating homo- or heteropolymers, respectively, that are structurally very diverse. Notably, EPSs—particularly in pathogens—are thought to be critical in host–microbe interactions, where they aid in adherence and colonization within the human host (8) and function in immunomodulation (9). Although very little is known about the function of bifidobacterial

EPS, it has been suggested to aid in tolerance of the bacterium to bile/acid (10), and has also been shown to serve as a growth substrate for elements of the gut microbiota (7).

We show that the commensal *Bifidobacterium breve* UCC2003 contains a bidirectional EPS-encoding genetic locus responsible for the production of a surface-attached EPS that provides resistance to both bile and acid *in vitro*. In mice, this surface EPS aids in long-term persistence and also mediates immune evasion, specifically in avoiding B-cell responses. Colonization of mice with EPS⁺, but not EPS[−] *B. breve*, also provides the host with protection after enteric pathogen challenge. This finding suggests that surface EPS on commensal bacteria can facilitate colonization of their host through evasion of potentially damaging immune responses, and in turn can provide direct health-promoting benefits, in this case via a reduction in pathogen colonization.

Results

Identifying and Characterizing an EPS Locus in *B. breve* UCC2003. The genome of *B. breve* UCC2003 (11) harbors a putative EPS-encoding locus (designated here as *eps*), which extends from Bbr_0430 to Bbr_451, and encompasses a 25.6-kb region that harbors 20 genes predicted to be involved in EPS biosynthesis (Fig. 1A and Table S1) and two transposase-encoding sequences (Bbr_0432 and Bbr_0433). The majority (i.e., 18 of 22) of these genes are organized as two adjacent but oppositely oriented gene sets, the first encompassing Bbr_0441 to Bbr_0434, designated here as the *eps1* operon, the second from Bbr_0442 to Bbr_451, designated as the *eps2* operon (Fig. 1A; see also below). The divergence in GC content (Table S1) suggests that the *eps* locus was acquired by horizontal gene transfer, as indicated also for other EPS-encoding loci (12).

Transcriptional analyses by promoter fusions, quantitative RT-PCR (qRT-PCR) and primer extension delineated the transcriptional units and promoter sequences of this locus, revealing that the gene encoding the only predicted priming glycosyl transferase, Bbr_0430, and a gene (Bbr_0431) putatively encoding a protein involved in EPS chain-length regulation, are each transcribed by a separate promoter and thus, based on their genetic location and orientation, monocistronic (Fig. 1A). Transcriptional analyses

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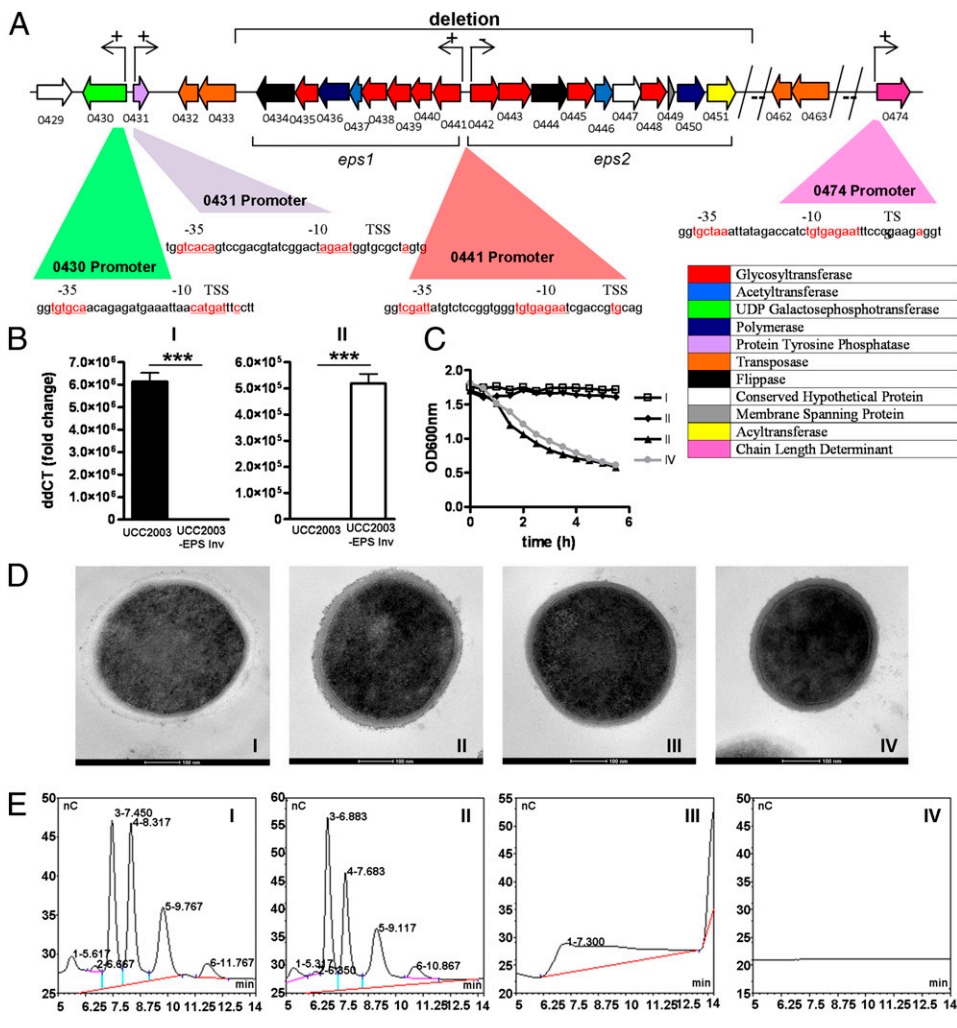


Fig. 1. Identifying and characterizing the EPS locus in *B. breve* UCC2003. (A) Schematic diagram of the *B. breve* UCC2003 EPS locus. For identification of promoters in the EPS: A "+" sign denotes a statistically significant ($P < 0.02$ by *t* test) difference in GUS activity between the generated promoter fusion constructs and the pNZ272 control in *B. breve* UCC2003 at the 12-h growth point, and a "-" sign denotes no statistically significant ($P < 0.02$ by *t* test) difference between the promoter fragment and the pNZ272 control in *B. breve* UCC2003 ($n = 3$). All cultures had similar growth rates. *eps1* and *eps2* refer to the two adjacent transcriptional units involved in EPS biosynthesis. // represents a number of genes between Bbr_0451 and Bbr_0462 or between Bbr_0463 and Bbr_0474. Transcriptional start sites, -10 and -35 sites of promoters are indicated; the color coding of the genes relates to their predicted function as indicated in the inset. (B) qRT-PCR analysis of the Bbr_0441 and Bbr_0442 transcriptional units. (I) Transcription of Bbr_0441 in strains UCC2003 and UCC2003-*-EPS Inv*; (II) transcription of Bbr_0442 in UCC2003 and UCC2003-*-EPS Inv*. (C) OD measurements (OD_{600nm}) of UCC2003 and its derivatives over a 5.5-h time period grown in batch culture without agitation; the observed drop in OD values for the *-EPS*⁻ derivatives is because of cell sedimentation. (D) Transmission electron microscopy images of *B. breve* UCC2003 (I) and isogenic derivatives UCC2003-*-EPSInv* (II), UCC2003-*-EPSdel* (III), UCC2003::Bbr_0430 (IV). (Scale bars, 100 nm.) (E) High performance anion-exchange chromatography with pulsed amperometric detection profiles of acid-hydrolyzed surface EPS isolated from UCC2003 and isogenic derivatives (see D for strain coding).

further showed that the *eps1* operon is constitutively transcribed in UCC2003 from a single promoter, but the *eps2* operon is transcriptionally silent (Fig. 1A and B). Serendipitously, a variant of UCC2003 was isolated, designated UCC2003-*-EPSInv*, which was found to exhibit constitutive transcription of the *eps2* operon, but *eps1* transcription was undetectable (Fig. 1B). Sequence analysis revealed that the observed transcriptional pattern of UCC2003-*-EPSInv* was the result of an inversion of a 282-bp DNA fragment, located within the intergenic region between *eps1* and *eps2*, thus causing promoter reorientation (Fig. 1A). Because this reoriented fragment is flanked on either side by a 75-bp sequence, which together form a near-perfect (1-bp mismatch) inverted repeat, this promoter reorientation is likely catalyzed by a site-specific DNA recombinase/invertase, reminiscent of the promoter switching phenomenon described for capsule synthesis modulation in *Bacteroides fragilis* (13).

To demonstrate that the *eps* locus is responsible for EPS/capsule production, an insertion mutant, designated UCC2003::Bbr_0430, was generated in the monocistronic Bbr_0430 gene, encoding the putative priming glycosyltransferase. In addition, a UCC2003 deletion derivative was fortuitously isolated (designated UCC-*-EPSdel*), which had lost the DNA region between two identical insertion sequence elements (from Bbr_0432/0433 to and including Bbr_0463/0464), which encompassed both *eps1* and *eps2* (Fig. 1A). Transmission electron microscopy showed that strains UCC2003 and UCC2003-*-EPSInv* produce an outer cell-surface layer, presumed to be a capsule consisting of surface EPS, which is absent in UCC2003-*-EPSdel* and UCC2003::Bbr_0430 (Fig. 1D) (these

strains are therefore designated as *-EPS*⁻). Interestingly, both *-EPS*⁻ strains were found to sediment during growth in liquid medium, but the *-EPS*⁺ strains remained in suspension (Fig. 1C). Surface EPS isolation and subsequent acid hydrolysis on the presumed *-EPS*⁻ and *-EPS*⁺ strains showed that no identifiable sugar peaks were present in the chromatograms obtained for the *-EPS*⁻ strains, whereas the chromatogram patterns observed for the *-EPS*⁺ strains revealed distinct monosaccharide profiles (Fig. 1E). This finding confirms that the two latter strains both produce an EPS surface layer, presumably each having a specific saccharide composition and structure. No detectable monosaccharide peaks were identified when we attempted to isolate EPS from spent growth medium in which *B. breve* UCC2003 had been cultivated, suggesting that little if any EPS is released from cells during growth in liquid medium.

***B. breve* UCC2003 Surface EPS Is Linked to Acid and Bile Resistance.** To investigate the potential relationship between UCC2003's ability to produce EPS and tolerance of low pH and bile salt-containing environments, growth profiles of UCC2003 and *-EPS*⁻ derivatives were monitored. When the two *-EPS*⁻ strains were grown at a pH of 5.0 or pH 4.0, they exhibited a significantly reduced ($P < 0.001$) growth rate and reached a lower end-point OD_{600nm}, compared with their *-EPS*⁺ counterparts (Fig. 2A). When these strains were grown in 0.3% bovine bile the *-EPS*⁻ strains exhibited significantly ($P < 0.01$) lower growth rates and lower end-point OD_{600nm} than the *-EPS*⁺ strain UCC2003 (Fig. 2A). These results show that the EPS layer has a protective effect under low pH and bile conditions.

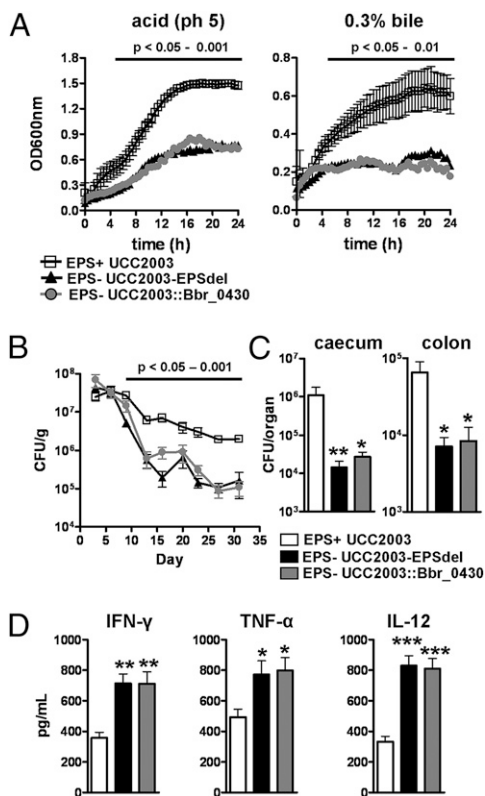


Fig. 2. *B. breve* surface EPS protects against acid and bile, facilitates in vivo persistence, and modulates cytokine expression from stimulated splenocytes. (A) Growth curve of UCC2003 (EPS⁺), UCC2003-EPSdel (EPS⁻), and UCC2003::Bbr_0430 (EPS⁻) in de Man-Rogosa-Sharpe (MRS) pH5.0 or in MRS supplemented with 0.3% bile over 24 h at 37 °C. Data represent mean \pm SD. (B) BALB/c mice were treated orally with $\sim 1 \times 10^9$ UCC2003, UCC2003-EPSdel, or UCC2003::Bbr_0430 on 3 consecutive days and bacterial numbers (CFU) in feces determined (data represent \log_{10} CFU/g feces \pm SD). (C) Organs were also removed on day 31 to determine CFU; columns show \log_{10} CFU/organ (\pm SD). (D) Spleens were harvested from naïve BALB/c mice and stimulated with wild-type (EPS⁺), deletion, and insertion mutants (EPS⁻) *B. breve* strains at 1:1 ratio for ~ 20 h. Cells were also stimulated with ConA (dotted line). Columns represent the mean \pm SD stimulation indices of splenocytes from 10 mice from two independent experiments. Significance was determined relative to mice treated with UCC2003 (EPS⁺) at the same time point using the Kruskal–Wallis test followed by Dunn’s multiple comparison test; * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$.

***B. breve* Surface EPS Impacts on Persistence, but Not Initial Colonization, in Mice.** To establish if bifidobacterial EPS production impacts on colonization ability and persistence patterns in BALB/c mice, viable counts of *B. breve* UCC2003 and various derivatives were determined from fecal samples over a 31-d period following a 3-d treatment period. First, a comparison was made between *B. breve* UCC2003 and *B. breve* UCC2003-EPSInv to establish whether expression of different EPS impacts on colonization or persistence of *B. breve*. No differences were observed between these two EPS⁺ strains in fecal counts. We next compared EPS⁺ *B. breve* UCC2003 with EPS⁻ *B. breve* strains and noted that from as early as 9 d posttreatment, and up until the end of the study, significantly lower ($P < 0.05$ – 0.001) bacterial numbers were recovered from mice treated with EPS⁻ strains compared with EPS⁺-treated animals (Fig. 2B). On day 31, the viable counts of the EPS⁻ strains within the murine caeca and colons were also significantly reduced ($P < 0.05$) compared with those of EPS⁺ strains (Fig. 2C).

***B. breve* Surface EPS Modulates Cytokine Levels in Stimulated Naïve Splenocytes.** To determine if *B. breve* EPS expression has an effect on host immune responses, we isolated splenocytes from naïve mice and stimulated them with isogenic EPS⁺ or EPS⁻ *B. breve* strains (Fig.

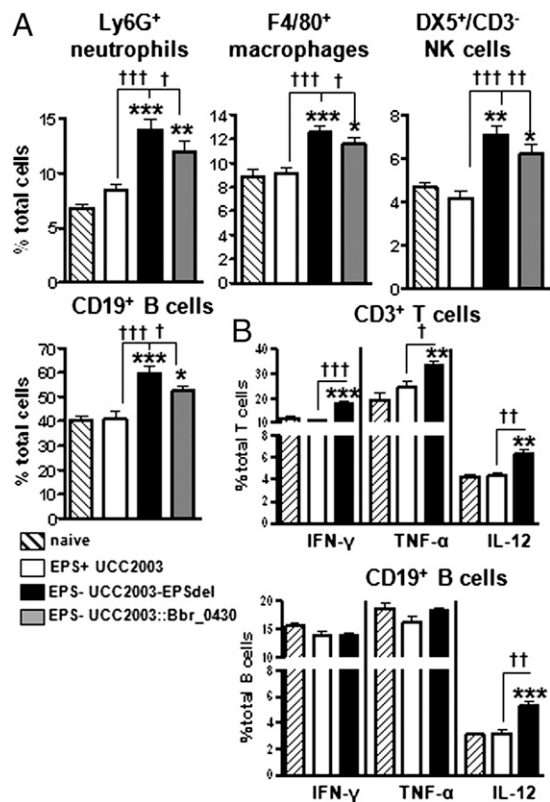


Fig. 3. *B. breve* surface EPS modulates recruitment and cytokine profile of immune cell populations in mice. (A) Cells were isolated from spleens of BALB/c mice 31 d after initial treatment, stained with fluorochrome-labeled mAb, and analyzed by flow cytometry. Columns represent the mean percentage \pm SD of at least eight mice from two independent experiments. (B) Isolated cells were stimulated for 6 h with BD Leukocyte Activation Mixture plus GolgiPlug in vitro, stained with surface mAb to determine CD3⁺ and CD19⁺ populations, and then permeabilized and stained with anticytokine fluorochrome-labeled mAb. Data represent percent of cytokine-positive cells out of total specific cell population \pm SD. * $P < 0.05$, ** $P < 0.01$, and *** $P < 0.001$ between naïve and *B. breve*-treated mice; † $P < 0.05$; †† $P < 0.01$, and ††† $P < 0.001$ between EPS⁺ and EPS⁻ using one-way ANOVA followed by Bonferroni’s multiple comparison test.

2D). All *B. breve* strains induced higher cytokine levels compared with unstimulated negative controls. However, cells stimulated with EPS⁺ had significantly ($P < 0.01$) lower levels of the proinflammatory cytokines IFN- γ , TNF- α , and IL-12 compared with UCC2003-EPSdel and UCC2003::Bbr_0430 (EPS⁻)-stimulated cells.

Treatment of Mice with EPS⁺ *B. breve* Causes Reduced Levels of Proinflammatory Immune Cells Compared with EPS⁻ Strains. To determine if murine treatment with EPS⁺ or EPS⁻ *B. breve* impacts on in vivo immune responses, spleens from untreated (naïve) mice or from mice given three oral doses of UCC2003 (EPS⁺), UCC2003-EPSdel (EPS⁻), or UCC2003::Bbr_0430 (EPS⁻) were examined by flow cytometry. Mice treated with UCC2003 (EPS⁺) did not exhibit any differences in either percentage or total cell number of any of the immune cells monitored compared with untreated mice. In contrast, mice treated with either of the EPS⁻ strains had significantly increased ($P < 0.01$) percentages and total cell numbers of Ly6G⁺ neutrophils, F4/80⁺ macrophages, DX5⁺/CD3⁻ NK cells, and CD19⁺ B cells compared with naïve mice (Fig. 3A and Table S2). Mice treated with EPS⁻ *B. breve* strains also had increased percentages and total immune cell numbers compared with EPS⁺-treated mice. Analysis of intracellular cytokine expression (Fig. 3B and Table S3) revealed that although the numbers of T cells did not change, their cytokine profile was significantly altered. Concentrating on the cytokines that were altered in the stimulated naïve splenocyte studies

(Fig. 2D), we observed that the percentage of IL-12, IFN- γ , and TNF- α -positive T cells, was significantly higher ($P < 0.01$) in mice treated with a EPS⁻ *B. breve* strain compared with both naïve or EPS⁺-treated mice. Analysis of B-cell populations revealed significant increases ($P < 0.01$) in the IL-12-producing subpopulation in EPS⁻-treated mice compared with naïve and UCC2003 (EPS⁺)-treated mice. Total cell-number examination revealed more IFN- γ - and TNF- α -producing B cells in mice treated with the EPS⁻ *B. breve* strain compared with either naïve or UCC2003-treated mice. Innate populations were also examined and demonstrated significantly lower ($P < 0.05$) proinflammatory cytokine profiles in EPS⁺ compared with EPS⁻ *B. breve*-treated mice (Fig. S1).

Absence of Surface EPS on *B. breve* UCC2003 Alters Its Antibody Responses. Mice treated with EPS⁻ *B. breve* strains had significantly higher ($P < 0.01$) total numbers of CD134⁺/CD19^{low}/B220^{low} plasma B cells compared with animals treated with the EPS⁺ strain or naïve mice on day 31 (Fig. 4A). Analysis of serum antibody titers revealed that although mice treated with EPS⁺ *B. breve* did mount a modest antigen-specific total Ig response, a much higher response was observed in mice treated with the EPS⁻ strain (Fig. 4B). This result was also the case for the serum antibody subtypes IgG3, IgG1, and IgG2a. Mice treated with the *B. breve* EPS⁺ strain had no detectable serum IgG3 titers above preimmune levels. Although overall fecal IgA titers were low, we again observed a significantly higher ($P < 0.01$) level in mice treated with the EPS⁻ *B. breve* strain compared with those treated with UCC2003.

We next wanted to determine if the presence of surface EPS would impact on agglutination with serum raised in mice treated with either EPS⁺ or EPS⁻ *B. breve* strains. From the images presented in Fig. 4C, *B. breve* UCC2003 (EPS⁺) was only weakly agglutinated with anti-UCC2003 serum, but not anti-EPS⁻ serum, but the EPS⁻ *B. breve* strain was shown to be strongly agglutinated with serum obtained from EPS⁻-treated mice, but not with anti-UCC2003 serum. Thus, it appears that EPS elicits only weak antibody responses, as well as potentially masking other *B. breve* UCC2003 surface antigens from exposure to antibody responses.

Efficient Colonization of *B. breve* Requires Evasion of B-Cell Responses. Given the apparent role of surface EPS in controlling host immune responses, we wanted to define the key adaptive host effector mechanism that is modulated by EPS. We focused on the role of B cells, because mice treated with EPS⁺ compared with EPS⁻ *B. breve* had reduced B-cell numbers together with diminished cytokine

and antibody profiles (Figs. 3 and 4). Treatment of B-cell-deficient mice (14) with UCC2003 (EPS⁺) led to similar bacterial numbers in the feces throughout the study compared with WT C57BL/6 controls (Fig. 4D). However, B-cell-deficient mice failed to reduce the number of EPS⁻ *B. breve* several weeks following colonization, which was observed in B-cell-proficient mice (Fig. 4D). We also observed this profile within the caeca at the end of the study (Fig. 4E). Colonization and immune responses in control C57BL/6 mice after treatment with isogenic EPS⁺ and EPS⁻ *B. breve* strains were similar to those observed in BALB/c animals. Taken together, these data indicate that *B. breve* surface EPS modulates B cells and is critical for long-term colonization.

Presence of Surface EPS on *B. breve* Directly Impacts on the Colonization and Persistence of the Gut Pathogen *Citrobacter rodentium*. To investigate if surface EPS impacts on host protection against an invading pathogen, mice were treated with *B. breve* EPS⁺ or EPS⁻ strains before oral gavage of bioluminescent *C. rodentium* (15). As early as 24 h postinfection, we observed significantly lower ($P < 0.05$) *C. rodentium* fecal counts from mice treated with EPS⁺ *B. breve* compared with EPS⁻ *B. breve*-treated mice, which continued throughout the study up until day 14 (Fig. 5A). By day 3, and up until the study end, we observed that EPS⁺ *B. breve*-treated mice had reduced numbers of *C. rodentium* compared with *C. rodentium*-infected mice not treated with *B. breve*. Whole-body bioluminescent data confirmed these observations (Fig. 5C). When organs were excised and imaged on days 6 and 8, we observed that mice treated with EPS⁻ *B. breve* and infected with *C. rodentium*, as well as untreated mice, showed strong luminescent signals from both the cecum and colon, but infected mice treated with EPS⁺ UCC2003 showed either undetectable or lower bioluminescence. At the end of the study (day 14), mice treated with EPS⁻ *B. breve* had detectable signals in both organs, with untreated mice no longer having any signal from the colon. Importantly, mice treated with EPS⁺ *B. breve* UCC2003 and infected with *C. rodentium* no longer had detectable luminescence in any of the organs (Fig. 5D), as also confirmed by viable count determination (Fig. 5B). Throughout the trial the colonization levels of the *B. breve* strains were similar to those described in our animal studies described above.

Discussion

There are numerous pathogenic roles described for EPS-producing bacteria. However, the precise biological role of EPS produced by commensal bacteria, such as bifidobacteria, has yet to be determined.

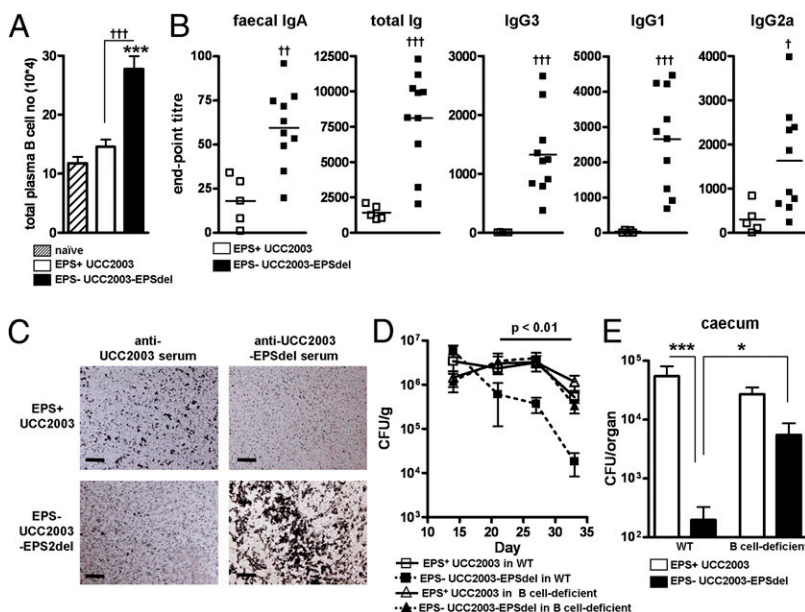


Fig. 4. *B. breve* EPS influences B-cell responses. (A) Cells were isolated from spleens of *B. breve*-treated mice on day 31 and data represent total cell number of CD19^{low}/B220^{low}/CD138^{high} plasma cells. Columns represent means \pm SD. Significant differences were determined by the one-way ANOVA followed by Bonferroni's multiple comparison test. *** $P < 0.001$. (B) Antibody titers were measured following treatment of BALB/c mice with UCC2003 (EPS⁺) and UCC2003-EPSdel (EPS⁻). Graphs show titers from individual serum and fecal samples (minus naïve titers) collected on day 31. Bars represent means from at least eight individual mice from two independent experiments. (C) Visualization of slide agglutination performed with the indicated antiserum and cultures of *B. breve*. Slides were counterstained with India ink and agglutination visualized by light microscopy (Original magnification, 20 \times). (Scale bars, 200 μ m.) (D) B-cell-deficient mice and C57BL/6 controls were treated orally with $\sim 1 \times 10^9$ EPS⁺ or EPS⁻ *B. breve* strains on 3 consecutive days and fecal CFU determined. (E) On day 14, organ counts were also determined. Data represent log₁₀ CFU (\pm SD) from eight individual mice. Significant differences determined by the Kruskal-Wallis test followed by Dunn's multiple comparison test. * $P < 0.05$; *** $P < 0.001$.

to the surface of bacteria may also prevent binding to host cells, and this may additionally explain why we observe the gradual reduction of EPS⁻ levels relative to levels of EPS⁺ *B. breve* in similarly treated mice. Using a B-cell-deficient mouse strain, we confirmed that *B. breve* UCC2003 surface EPS is crucial for persistence within the murine host, through subversion of B-cell responses, similar to that used by many pathogenic bacteria (20). However, as far as we are aware, this study is unique in looking at EPS-dependent modulation of B-cell responses in a commensal bacterium.

Using the murine pathogen *C. rodentium* as a model for human enteropathogenic *Escherichia coli* and enterohemorrhagic *E. coli* (which also occupy the same environmental niche as *B. breve*) (21), we found that surface EPS expression had a profound effect on initial pathogen colonization, as well as burden. EPS⁺ and EPS⁻ *B. breve* had similar colonization levels at early time points. Therefore, the mechanism by which UCC2003 reduces *C. rodentium* levels does appear to be solely because of total *B. breve* load. One of the mechanisms by which UCC2003 may “curb” pathogen levels within its host, compared with EPS⁻ derivatives, may be through the ability to form biofilms. Furthermore, the absence of surface EPS may lead to this otherwise immunological silent commensal being detected as a foreign bacterium, and therefore stimulating a strong immune response. This finding may explain the significantly higher levels of *C. rodentium* observed in EPS⁻ treated mice, compared with *C. rodentium*-infected mice not treated with *B. breve*, as coinfection can result in increased pathogen burden and associated pathology (22). More studies are required to properly dissect this protective mechanism; however, this surface EPS-dependent pathogen defense represents an exciting new avenue for probiotic research.

In conclusion, we have comprehensively characterized the biological functions of surface EPS in the human commensal *B. breve* UCC2003. Our findings are consistent with an EPS-mediated strategy that prevents a commensal bacterium from evoking a strong adaptive immune response within the local environment. Furthermore, surface EPS production by UCC2003 provides protection against infection of a murine pathogen, thus representing a mode of action to elicit positive health benefits.

Materials and Methods

Bacterial Techniques. For detailed information on bacterial strains, culturing conditions, and analysis techniques used in this study see *SI Materials and Methods* and *Table S4*.

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Animals and Colonization by *B. breve*. Male and female BALB/c mice (6–8 wk old) were purchased from Harlan. Both wild-type and B-cell-deficient B6.129S2-*Igh-6^{tm1Cgn}/J* C57BL/6 mice, were obtained from the Jackson Laboratory. Animal husbandry and experimental procedures were approved by the University College Cork ethics committee. Groups of mice were orally gavaged with $\sim 1 \times 10^9$ CFU per mouse for 3 consecutive days with *B. breve* strains. Fresh fecal samples were plated at several time points for 31 d posttreatment. At the end of the studies mice were killed, and small intestine, cecum, and colons were excised and organ homogenates plated.

Flow Cytometry. Single-cell suspensions were prepared and stained as previously described (23), with monoclonal antibodies laid out in *Table S5*. For intracellular staining, cells were incubated with BD Leukocyte Activation Mixture with BD GolgiPlug or GolgiPlug alone (BD Biosciences) for 6 h. Samples were acquired on a FACSLSR II and data were analyzed using DIVA software.

Evaluation of Antibody Responses. Serum samples from mice were obtained at the end of the study (day 31) and analyzed for the presence of total Ig, IgG1, IgG2a, and IgG3, as previously described (23) and in *SI Materials and Methods*. For agglutination assays, *B. breve* suspensions were incubated with appropriate serum and visualized with India ink at a magnification of 20 \times using an Olympus microscope.

***C. rodentium* Infections and Bioluminescent Imaging of Mice.** Groups of BALB/c mice were treated with the appropriate *B. breve* strains as described above. A group of untreated mice was also included as a control. *C. rodentium* was grown and administered to mice. At selected time points postinfection, assessment of bioluminescence from living animals was measured by the use of an IVIS50 system (Xenogen), as previously described (15) and in *SI Materials and Methods*.

Statistical Analysis. Experimental results were plotted and analyzed for statistical significance with Prism4 software (GraphPad) using Student *t* test, one-way ANOVA followed by Bonferroni post hoc correction and ANOVA Kruskal–Wallis test with Dunn’s multiple comparison test. A *P* value of < 0.05 was used as significant in all cases.

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