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## BVES Is Required for Maintenance of Colonic Epithelial Integrity in Experimental Colitis by Modifying Intestinal Permeability

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### Abstract

Blood vessel epicardial substance (BVES), or POPDC1, is a tight junction-associated transmembrane protein which modulates epithelial to mesenchymal transition (EMT) via junctional signaling pathways. There have been no *in vivo* studies investigating the role of BVES in colitis. We hypothesized that BVES is critical to maintaining colonic epithelial integrity. At baseline, *Bves*<sup>-/-</sup> mouse colons demonstrate increased crypt height, elevated proliferation, decreased apoptosis, altered intestinal lineage allocation, and dysregulation of tight junctions with functional deficits in permeability and altered intestinal immunity. *Bves*<sup>-/-</sup> mice inoculated with *Citrobacter rodentium* had greater colonic injury, increased colonic and mesenteric lymph node bacterial colonization, and altered immune responses after infection. We propose that increased bacterial colonization and translocation result in amplified immune responses and worsened injury. Similarly, dextran sodium sulfate (DSS) treatment resulted in greater histologic injury in *Bves*<sup>-/-</sup> mice. Two different human cell lines (Caco2 and HEK293Ts) co-cultured with enteropathogenic

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*E. coli* showed increased attaching/effacing lesions in the absence of BVES. Finally, *BVES* mRNA levels were reduced in human ulcerative colitis (UC) biopsy specimens. Collectively, these studies suggest that BVES plays a protective role both in ulcerative and infectious colitis and identify BVES as a critical protector of colonic mucosal integrity.

## Keywords

BVES; inflammatory bowel disease; epithelial injury; colitis

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## Introduction

Inflammatory Bowel Disease (IBD) affects over 1.6 million people in the United States, causing numerous and debilitating complications such as malnutrition, infection, pain, and an elevated risk of colorectal cancer.<sup>1,2</sup> The fundamental pathophysiology of IBD is not well understood. Intestinal mucosal homeostasis relies on continual proliferation, differentiation, and repair of the epithelium, and these processes are challenged in response to injury.<sup>3-5</sup> Maintenance of intestinal integrity requires competent epithelial barrier function as well as intact repair programs to respond to intestinal challenges.<sup>3-5</sup> The epithelial barrier is “leaky” in patients with IBD; multiple studies demonstrate lower epithelial resistance and increased permeability of both inflamed and non-inflamed mucosa in Crohn’s disease (CD) and ulcerative colitis (UC).<sup>6,7</sup> Tight junctions are apical complexes critical to the maintenance of epithelial barriers.<sup>8,9</sup> Disruption of tight-junctional complexes is a key contributing factor to epithelial barrier dysfunction.<sup>6-7,10</sup> For example, tight junction-associated proteins, such as F11r, which we will refer to as junctional adhesion molecule-A (JAM-A), and MYH2, which we will refer to as non-muscle myosin II (NMII), are protective in the dextran sodium sulfate (DSS) model of experimental colitis.<sup>11-12</sup> Loss of either JAM-A<sup>11</sup> or NMII<sup>12</sup> resulted in low grade inflammation and increased intestinal permeability, but interestingly, the mice were able to maintain intestinal homeostasis and were otherwise normal at baseline. However, upon colitogenic challenge, these mice were unable to compensate, resulting in markedly more severe injury.

Blood vessel epicardial substance (BVES) is a tight junction-associated, three-pass transmembrane protein originally discovered from a cDNA screen of the developing heart.<sup>13,14</sup> BVES is expressed in epithelial tissues, including the intestinal epithelium, and regulates epithelial-to-mesenchymal transition (EMT).<sup>13,15-20</sup> We have previously demonstrated that BVES regulates stem cell programs in the small intestine,<sup>20</sup> colonic epithelial phenotypes *in vitro*,<sup>15</sup> and can promote inflammatory tumorigenesis *in vivo*,<sup>21</sup> but its role in colitis remains unexplored. We hypothesized that BVES is critical to maintaining epithelial integrity after colonic injury.

At baseline, *Bves*<sup>-/-</sup> mice exhibited higher colonocyte proliferation, decreased apoptosis, defects in intestinal secretory lineage allocation, and dysregulation of tight junction proteins. *Ex vivo* studies, electron microscopy (EM), and 2-D colonoid cultures demonstrated increased permeability in *Bves*<sup>-/-</sup> colons. Moreover, *Bves*<sup>-/-</sup> mice displayed dysregulated mucosal chemokine and cytokine production. After inoculation with *Citrobacter rodentium*,

*Bves*<sup>-/-</sup> mice had exacerbated colitis along with increased colonic and mesenteric lymph node bacterial colonization, decreased proliferation and stem cell marker expression, and skewing of mucosal chemokines and cytokines. In confirmatory studies, *Bves*<sup>-/-</sup> mice treated with DSS also had increased histologic injury in both acute and injury-repair models. *BVES* silencing in HEK293T and Caco2 cells resulted in increased attaching/effacing lesions and *CXCL8* mRNA expression after EPEC infection, while HEK293T cells overexpressing *BVES* had fewer attaching/effacing lesions and decreased mRNA expression of *CXCL8*. Lastly, analysis of human UC mucosal biopsies demonstrated decreased *BVES* expression, suggesting that its downregulation is relevant to intestinal injury processes. Collectively, these findings indicate that BVES regulates the mouse colonic microenvironment, protecting against intestinal epithelial injury and inflammation, and may play a protective role in IBD.

## MATERIAL AND METHODS

### Mouse Models

Wildtype (WT) (C57BL/6 background) were obtained from the Jackson Laboratories. *Bves*<sup>-/-</sup> mice have previously been described in detail.<sup>15</sup> All experiments were performed with 8 to 12 week old male and female mice on C57BL/6 background under guidelines approved by the Vanderbilt University Institutional Animal Care and Use Committee.

### Functional Permeability Analysis

These experiments were done with as previously described.<sup>22</sup> Briefly, sections of distal colon from eight to twelve-week-old male mice were mounted between the two halves of the Ussing Chamber (1.12 cm<sup>2</sup> opening). The mucosal and serosal sides of chambers were connected to circulating reservoirs which contained Ringer's solution with 5 mM mannitol. A gas lift column of 95% O<sub>2</sub> maintained oxygenation and pH 7.4 of reservoirs. *I*<sub>sc</sub> was continuously monitored on a strip chart recorder, and electrical resistance was calculated using Ohm's law. After baseline potential difference and resistance was measured, a fluorescent tracer (FITC-dextran, 4 kDA, #46944, Sigma) at a concentration of 80 mg/ml was placed on one side (luminal bath) and concentration was measured three hours after beginning experiment on the other side (serosal bath) via spectrophotofluorometry, with an excitation wavelength of 492 nm and emission wavelength of 518 nm.

### Electron Microscopy

Specimens were processed for transmission electron microscopy (TEM) and imaged in the Vanderbilt Cell Imaging Shared Resource-Research Electron Microscopy facility.

**Embedding**—Samples were fixed in 2.5% gluteraldehyde in 0.1M cacodylate buffer, pH7.4 at room temperature (RT) 1 hour then transferred to 4°C, overnight. The samples were washed in 0.1M cacodylate buffer, then incubated 1 hour in 1% osmium tetroxide at RT, and then washed with 0.1M cacodylate buffer. Subsequently, the samples were dehydrated through a graded ethanol series and then 3 exchanges of 100% ethanol. Next, the samples were incubated for 5 minutes in 100% ethanol and propylene oxide (PO) followed by 2 exchanges of pure PO. Samples were then infiltrated with 25% Epon 812 resin and

75% PO for 30 minutes at RT. Next, they were infiltrated with Epon 812 resin and PO [1:1] for 1 hour at RT then overnight at RT. The next day the samples went through a [3:1] (resin: PO) exchange for 3–4 hours and were incubated with pure epoxy resin overnight. Samples were then incubated in two more changes of pure epoxy resin and allowed to polymerize at 60°C for 48 hours.

**Sectioning and Imaging**—500–1000 nm thick sections were cut for ultra-structure identification. Then 70–80 nm ultra-thin sections were cut from the region of interest and collected on 300-mesh copper grids and post-stained with 2% uranyl acetate and then with Reynold's lead citrate. Samples were subsequently imaged on the Philips/FEI Tecnai T12 electron microscope at various magnifications.

### Colonoid Transepithelial Resistance (TER) measurements

Colonoid cultures were derived according to previously published protocols.<sup>23</sup> Briefly, the colon was dissected, flushed with ice cold phosphate buffered saline (PBS), and opened lengthwise. Colons were cut into 5 mm pieces before being transferred to chelation buffer (2 mM EDTA in Dulbecco's PBS) and rocked for 90 minutes at 4°C. After removing the chelation buffer, cold shaking buffer (PBS with 43.3 mM sucrose and 54.9 mM sorbitol) was added and tissues were gently shaken for 4 minutes to free colonic crypts. Colonic crypts were plated in growth factor reduced Matrigel (Corning, 356231) at a concentration of 25 crypts per 5  $\mu$ L plug. After polymerization, Matrigel was overlaid with 500  $\mu$ L mini-gut media (Advanced DMEM/F12, B27, N2, Pen-Strep, L-glutamine, HEPES) and supplemented with EGF (R&D systems, 2028-EG-200), Noggin (R&D Systems, 1967-NG-025/CF) and R-spondin (Vanderbilt Antibody and Protein Resource). For transwell plating, colonoids were collected and dissociated using TrypLE Express (Gibco, Life Technologies, 12604-013). Following dissociation, cells were plated on 0.4 $\mu$ m transwell filters (Corning, 3413) coated with Collagen 1 (Gibco, A10483-01) and cultured in mini-gut media supplemented with 10 $\mu$ M Y27632 (R&D Systems, 1254/10). After 1 day, mini-gut media was replaced by Advanced DMEM/F12 containing 1% penicillin/streptomycin, 1X Glutamax (Gibco, Life Technologies, 35050-061), and 10% FBS. Trans-epithelial resistance (TER) was measured by Ohm meter daily for 5 days.

### Western Blot Analysis

For western blot analysis, crypts were isolated by the same method for colonoid culture described above. Crypts were pelleted by centrifugation and resuspended in RIPA buffer, phosphatase inhibitor cocktail 2 (Sigma, P5726), phosphatase inhibitor cocktail 3 (Sigma, P0044), protease inhibitor (Sigma, P8340), and 2X Laemmli buffer and then boiled for 5 minutes before analysis by SDS-PAGE electrophoresis. Membranes were blocked using Odyssey blocking buffer for 30 minutes and then blotted with anti E-cadherin (BD Transductions, #610182), anti-ZO-1 (Life Technologies, #617300), anti-Claudin7 (Invitrogen #QL221418), anti-GAPDH (Cell Signaling, #5174S), anti- $\beta$ -actin (Sigma, #116M4801V), anti-myosin light chain 2 (Cell Signaling, #8505S), anti-phospho-myosin light chain 2 (Cell Signaling, #3674S), or Myosin 2C (Cell Signaling, #8189S). All antibodies were used at 1:1000 concentration (except anti- $\beta$ -actin, 1:2000) in Odyssey blocking buffer with 0.1% Tween-20 and incubated overnight at 4°C. LiCor secondary

antibodies were used at 1:10,000 dilution and incubated for 2 hours at room temperature. Quantification of western blot band intensity was conducted using LiCor Image Studio.

### Induction of *Citrobacter rodentium* colitis

Mice were orally inoculated with *C. rodentium* as done previously.<sup>24–27</sup> Bacteria were grown overnight in Luria broth (LB), and subculture was performed the following day to ensure live bacteria were in exponential phase. Mice were infected by oral gavage with 0.1 mL of broth containing  $5 \times 10^8$  CFU of *C. rodentium*. Subculture was performed the following day to ensure live bacteria were in exponential phase. Control mice received sterile broth. At 14 days post-inoculation, the animals were sacrificed and the colons were removed, cleaned, and Swiss-rolled for histology. For colonization studies, distal colon pieces or mesenteric lymph nodes were homogenized, serially diluted in LB, and plated on MacConkey agar plates. For colonic tissues, colonies were counted after 24 hours and CFU/g was calculated. For mesenteric lymph nodes, colonies were counted after 48 hours and CFU/g was calculated. The *C. rodentium* histology slides were reviewed and scored in a blinded fashion by a gastrointestinal pathologist (M. B. P.). Both acute (neutrophilic) and chronic (lymphocytic) inflammation, extent of inflammation, and epithelial damage were each scored on 0–3 scale.<sup>24,27</sup> The aggregate histologic injury score was the sum of acute and chronic inflammation multiplied by the extent of inflammation, plus the epithelial injury (0–21).<sup>27</sup>

### Immunohistochemistry

For histological assessment, colons were removed, Swiss-rolled, and fixed in 10% formalin overnight prior to submission to the Vanderbilt University Translational Pathology Shared Resource (TPSR) core for processing and paraffin embedding. For immunohistochemistry (IHC), 5  $\mu$ m sections were cut, dewaxed, hydrated, and endogenous peroxidase activity quenched with 0.03% hydrogen peroxide in MeOH.<sup>28–30</sup> Antigen Unmasking Reagent (Vector Laboratories, H-3300) was used for antigen retrieval. The Vectastain ABC Elite System (Vector Laboratories, AK-5002) was then used according to manufacturer's instructions. Primary antibodies were added overnight at 4°C with isotype-matched antibodies (negative controls) on serial sections. The following antibodies were used: anti-phospho-Histone H3 (pH3) Ser10 antibody (1:150, #2465253, Millipore), anti-CgA (1:1000, #280170, ImmunoStar Inc.), anti-CD3 antibody (1:200, MCA500GA/1110, BioRad). Periodic Acid Schiff (PAS) staining for goblet cells and hematoxylin and eosin (H&E) staining was performed by the TPSR core. Apoptotic cells were identified using the ApopTag Plus Peroxidase In Situ Apoptosis Kit (Millipore, S7101) according to the manufacturer's protocol. Quantification was generated by counting cells in 100 sequential, well-aligned crypts from the distal colon and presented as the mean number of positive cells per crypt  $\pm$  standard error (SE).

For fluorescent IHC on paraffin-embedded sections, antigen retrieval consisted of 10 minute boiling in 10mM sodium citrate (pH 6). After blocking, samples were incubated with the following primary antibodies overnight at 4°C: anti-Ly6B.2 antibody (1:100, BioRad, MCA771G), anti-B220 antibody (1:500, BD Pharmingen, 550286), anti-E-cadherin (1:500, #610182, BD Transduction Laboratories), or anti-ZO-1 (1:500, Life Technologies,

#617300). Secondary antibodies were conjugated to either 488 or 594 Alexafluor dyes (1:1000, Invitrogen) and incubated for 2 hours at room temperature. Nuclear staining done with ProLong Gold antifade reagent with DAPI (Invitrogen, 1846939).

For fluorescent IHC on frozen sections, colons were removed, Swiss-rolled, and immediately frozen in Optimal Cutting Temperature (OCT) compound (Tissue-Tek, #4583). Five  $\mu\text{m}$  sections of OCT embedded tissue were cut using a cryostat. Sections were fixed with 1% paraformaldehyde, permeabilized with Triton X-100, and blocked/quenched with 10% normal goat serum (50% mM  $\text{HN}_4\text{Cl}$ -filtered). Samples were then incubated with the following primary antibodies overnight at 4°C: anti-phospho-myosin light chain 2 (1:200, Cell Signaling, 3674S), and Myosin IIc (1:200, Cell Signaling, 8189S), Myosin IIb (1:200, Cell Signaling, 8824T), and Phalloidin (1:500, Biotium, 00043T). Secondary antibodies were conjugated to either 488 or 594 Alexafluor dyes (1:1000, Invitrogen) and incubated for 2 hours at room temperature. Nuclear staining done with ProLong Gold antifade reagent with DAPI (Invitrogen, 1846939).

### qRT-PCR analysis

RNA from *Bves*<sup>-/-</sup> or WT colons was isolated using the Direct-zol RNA MiniPrep Kit (Genesee Scientific, #11–311) or RNeasy Mini Kit (Qiagen, #74106) and cDNA was synthesized using the qScript cDNA synthesis kit (Quantabio, #95047-500) or iScript cDNA synthesis kit (Bio-rad, #1708891) from 1  $\mu\text{g}$  of total RNA. 1  $\mu\text{L}$  of cDNA was used as a template in each subsequent PCR reaction. Taqman probes were purchased from Applied Biosystems and *CXCL8/IL-8* (Hs00174103) and *BVES* (Hs00362584) were analyzed using the delta-delta Ct method and normalized to *GAPDH* (Hs02758991).

### Chemokine and Cytokine profiling

Fresh excised colon was rinsed and homogenized in CelLytic MT Mammalian Tissue Lysis/Extraction buffer (Sigma, St Louis, MO). Colonic tissue lysates were analyzed using the 25-analyte MILLIPLEX MAP Mouse Cytokine/Chemokine Magnetic Bead Panel (EMD Millipore) on a FLEXMAP 3D instrument. Protein concentration was measured using the Bio-Rad DC protein assay kit (Bio-Rad Laboratories). Concentration of each chemokine or cytokine was presented as pg/mg colon tissue protein.

### Dextran Sodium Sulfate (DSS) treatment

**Acute DSS injury**—Eight to twelve-week-old WT and *Bves*<sup>-/-</sup> C57BL/6 mice were treated with 4% DSS (Affymetrix, #14489) in the drinking water for 5 days and then sacrificed. Colonoscopy (Karl Storz, Veterinary Endoscopy) was performed before sacrifice and videos were scored using the murine endoscopic index of colitis severity (MEICS),<sup>31</sup> which is a 15-point score based on thickening of the colon (0–3), changes of the vascular pattern (0–3), fibrin visibility (0–3), granularity of the mucosal surface (0–3), and stool consistency (0–3). After sacrifice, colons were Swiss-rolled and analyzed for histologic injury in a blinded fashion by a gastrointestinal pathologist (M.K.W.) via calculation of an ulceration score, as previously described.<sup>32</sup> Briefly, the scoring system utilized a score of 0–4 to grade the extent of destruction of the crypt architecture. A score of 0 indicated normal crypt architecture, 1 indicated effacement of the bottom third of the crypt, 2 indicated



effacement of the bottom 2/3 of the crypt, 3 indicated complete effacement of the crypt, and 4 indicated complete crypt architectural effacement along with an absence of the normal epithelial surface.

**DSS injury repair model**—Eight to twelve-week-old WT and *Bves*<sup>-/-</sup> C57BL/6 mice were treated with 4% DSS for 5 days, allowed to recover for 5 days, and then sacrificed. As in the acute DSS model, colons were Swiss-rolled and histologic injury calculated based upon ulceration score.

**Chronic DSS model**—Eight to twelve-week-old WT and *Bves*<sup>-/-</sup> C57BL/6 mice were treated with 4% DSS for 5 days and then allowed to recover for 10 days. This was repeated two additional times. Again, colons were Swiss-rolled and histologic injury calculated as described above.

### Cell Culture

HEK293T and Caco2 cells were grown in DMEM containing 10% FBS, 2 mM glutamine, 100 U/ml penicillin, and 100 µg/ml streptomycin. Polyethylenimine (PEI) at a concentration of 1 µg/mL was used for all transfection experiments. An empty vector was used to ensure equal quantities of cDNA were transfected. *BVES* was knocked down using siRNA (Santa Cruz, #60295). siRNA was transfected using Lipofectamine RNAiMAX (Invitrogen, #13778-150). Scrambled siRNA (Santa Cruz, #37007) was used as control.

### Plasmids

BVES expression plasmids were generated using Gateway cloning (Invitrogen). pENTR vectors containing BVES (HsCD00368575) were shuttled into pcDNA 3.1V5 using IR clonase (Invitrogen).

### EPEC infection/FAS test

EPEC strain E2348/69 was maintained on Luria-Bertani agar plates. The bacterial concentration was estimated to be  $1 \times 10^8$  bacteria/ml per OD unit at 600 nm, as calculated by plating. HEK293T cells were plated on coverslips and infected with EPEC at a multiplicity of infection (MOI) of 10 for 3 hours. Cells were washed with PBS, fixed with 3% paraformaldehyde, permeabilized in 0.1% Triton X, and stained with an actin probe (ActinGreen 488, Molecular Probes, #1608515) for 30 minutes. Coverslips were mounted on glass slides with ProLong Gold antifade reagent with DAPI (Invitrogen, 1846939) and visualized with a Zeiss LSM880 confocal microscope. 15–20 photos of high-powered fields (63× or 80×) were taken and the number of attaching/effacing lesions were quantified by a blinded observer.

### RNA Scope

High-resolution RNA *in situ* hybridization was performed using the RNAscope® 2.5 HD Detection Kit – BROWN according to the manufacturer's protocol (ACDBio, document #322452 and #322310). Antigen retrieval was performed under standard pretreatment conditions as specified by the manufacturer. Probes were directed against BVES (catalog

#434599), MsPPIB (positive control, catalog #313911), and DapB (negative control, catalog #310043).

## Human Subjects

Tissues were obtained during colonoscopies at the Vanderbilt University Medical Center, following a protocol approved by the Vanderbilt Institutional Review Board.<sup>28</sup>

## Statistical Methods

Analyses comparing two groups were analyzed using the Mann-Whitney non-parametric test. Kruskal-Wallis was used to compare multiple groups. Survival curve analysis was done using the Log-rank (Mantel-Cox) test. Data are presented as the mean  $\pm$  the standard error of the mean (SEM) in bar graphs and a line identifying the mean is shown when all data points are plotted. All of these analyses were performed using GraphPad Prism®7.0c (San Diego, CA, USA).  $P < 0.05$  was considered statistically significant.

## RESULTS

### BVES regulates colonic crypt homeostasis

While it is known that loss of BVES increases susceptibility to colitis associated carcinoma<sup>21</sup> and regulates EMT,<sup>15</sup> a process critical in injury repair programs of the intestine,<sup>33,34</sup> the impact of its deletion on colonic homeostasis *in vivo* has not previously been examined. *Bves* messenger RNA is detectable in the colonic epithelium and is distributed uniformly from the base to luminal surface of the crypt in WT mice (Supplementary Figure 1). To determine if BVES deletion alters colonic phenotypes, we histologically characterized colons obtained from *Bves*<sup>-/-</sup> mice. Morphometric analysis demonstrated that *Bves*<sup>-/-</sup> mice had significantly greater crypt height (Figure 1A). In determining the basis for this phenotype, we noted via TUNEL staining that crypt apoptosis was decreased (Figure 1B) and phospho-histone H3 IHC revealed that proliferation was augmented in *Bves*<sup>-/-</sup> mice (Figure 1C), suggesting an imbalance in crypt growth/clearance kinetics which may contribute to the thickened mucosa observed in these mice. Abnormalities were not limited to growth parameters, as lineage allocation analysis revealed that the numbers of PAS-labeled goblet cells (Figure 1D) and Cga-labeled enteroendocrine cells (Figure 1E) were increased as compared to WT mice. Thus, BVES regulates colonic proliferation, apoptosis, lineage allocation, and crypt morphology, indicating a previously unrecognized role for BVES in influencing intestinal homeostasis.

### BVES contributes to epithelial barrier function

As we have previously identified that knocking down BVES with either siRNA or dominant negative approaches led to epithelial junctional compromise *in vitro*,<sup>15</sup> we hypothesized that the *Bves*<sup>-/-</sup> colon might also have abnormalities in tight junction-associated phenotypes such as permeability. While ultrastructural analysis of tight junctional complexes in *Bves*<sup>-/-</sup> colonic epithelium was unremarkable, measurements of intercellular lateral membrane boundaries demonstrated dilated intercellular spaces in *Bves*<sup>-/-</sup> colons as compared with WT (Figure 2A). Consistent with this observation, *Bves*<sup>-/-</sup> *ex vivo* cultures maintained in Ussing chambers demonstrated increased paracellular movement of FITC-Dextran



complexes after three hours, indicating barrier incompetency (Figure 2B). Finally, 3D *Bves*<sup>-/-</sup> colon organoids displayed reduced transepithelial resistance when grown on transwells as a 2D monolayer (Figure 2C). Together, these data indicate that the epithelial barrier is compromised in *Bves*<sup>-/-</sup> mice.

We next examined tight junction composition of the colonic crypts using a candidate gene approach. Immunoblot for Claudin 7 (Figure 2D) showed a significant decrease in protein levels. However, there was a trend towards increase in ZO-1 protein expression. We then performed qPCR for several other tight junction proteins (Figure 2F). There was a significant increase in *Claudin2*, *JAM-A*, and *Zo-1* mRNA expression. There was no change in mRNA expression in *Claudin1*, *Claudin3*, or *Claudin11*. Given that BVES has previously been shown to regulate RhoA signaling in the LIM2405 CRC cell-line,<sup>15</sup> we immunoblotted for phosphomyosin light chain 2 (pMLC) and myosin light chain 2 (MLC). While the immunoblot for pMLC and total MLC did not show a difference between WT and *Bves*<sup>-/-</sup> colonic crypt isolates, immunofluorescence of the epithelium shows an increase in pMLC staining on the luminal side of the crypt (Figure 2E). We then compared adherens junction proteins E-cadherin and  $\beta$ -catenin expression in colonic crypts by immunoblot and/or immunofluorescence and saw no difference in levels or localization (Supplementary Figure 2A–B). Finally, immunoblot showed a significant increase in myosin 2C expression, while immunofluorescence for phalloidin, myosin 2B, and myosin 2C showed no difference in localization (Supplementary Figure 3A–C). Thus, BVES deletion leads to dysregulation of several tight junction associated proteins in the colon, with the net effect being increased permeability.

### ***Bves*<sup>-/-</sup> mice display increased lymphoid aggregates at baseline**

Since *Bves*<sup>-/-</sup> mice exhibit increased permeability, we hypothesized that they would have alterations in immune cells within the colon. We first observed that there were increased lymphoid aggregates in the colonic mucosa in *Bves*<sup>-/-</sup> mice (Figure 3A) with an increase in fraction of T cells and myeloid cells (Figure 3B–C) and no detectable difference in B cells within these lymphoid aggregates (Figure 3D).

### **BVES maintains colonic epithelial integrity after *C. rodentium* colitis**

In order to analyze whether the aforementioned morphometric changes (i.e crypt hyperplasia), increased permeability, and increased lymphoid aggregate population increased the susceptibility of *Bves*<sup>-/-</sup> mice to develop experimental colitis, we inoculated mice with *C. rodentium*, which is a widely used model to induce colonic epithelial injury and is useful in interrogating intestinal repair responses. Both WT and *Bves*<sup>-/-</sup> mice lost weight after *C. rodentium* inoculation; however, by day 3 post-inoculation, *Bves*<sup>-/-</sup> mice trended towards increased weight loss from day 4 to day 14 (Figure 4A). At necropsy, the colon weight as a percent of body weight as well as colon weight/colon length was significantly greater in the *Bves*<sup>-/-</sup> mice (Figure 4B–C). Histological review of H&E-stained Swiss-rolled colons (Figure 4D) by a gastrointestinal pathologist examining metrics of epithelial ulceration, inflammatory infiltrate, submucosal edema, and impaired regeneration revealed worse injury in *Bves*<sup>-/-</sup> colons (Figure 4E). *Bves*<sup>-/-</sup> mice also demonstrated increased colonization of the

colonic mucosa (Figure 4F) and mesenteric lymph nodes (Figure 4G) after *C. rodentium* infection.

One consequence of increased bacterial colonization may be that WT mice have a greater proliferative response than *Bves*<sup>-/-</sup> colons in response to *C. rodentium*. In other words, the increase in proliferation from pre-injury to post-injury is greater in WT mice (Figure 4H). This is important because increased epithelial proliferation must occur to efficiently reconstitute epithelial barrier properties. We hypothesized that an increased stem cell compartment could explain this change in proliferation, as an increased stem cell number would result in an increased number of proliferating cells. As such, we compared mRNA expression of stem cell markers from colon epithelial isolates both before and after *C. rodentium* infection. At baseline, *cMyc* was significantly upregulated in *Bves*<sup>-/-</sup> colon epithelium while there was a trend towards an increase in stem cell markers *Lgr5* and *Lrig1* (Figure 4I). After *C. rodentium* infection, *cMyc* and *Lgr5* were both upregulated in WT while they were both down-regulated in *Bves*<sup>-/-</sup> mice relative to baseline. Though not statistically significant, *Lrig1* was upregulated after *C. rodentium* in both WT and *Bves*<sup>-/-</sup> mice but to a lesser degree in *Bves*<sup>-/-</sup> mice (Figure 4J). Finally, we found no differences in apoptosis (Supplementary Figure 4) between cohorts. As such, BVES loss predisposes mice to increased bacterial colonization (of both the colon and mesenteric lymph nodes) and thus more severe colitis. Moreover, *Bves*<sup>-/-</sup> mice have impaired injury repair programs, as evidenced by its relative hypoproliferative response and compromised stem cell compartment.

### ***Bves*<sup>-/-</sup> mice demonstrate altered immune responses at baseline and after *C. rodentium* infection**

Since *C. rodentium* uncovered increased histologic injury and impaired injury repair programs, we hypothesized that *Bves*<sup>-/-</sup> mice would also have alterations in immune response. Tissue specific chemokine and cytokine production was evaluated in *Bves*<sup>-/-</sup> colons using Luminex profiling. At baseline, there was increased protein expression of CCL5, IL-5, IL-15, and IL-1 $\alpha$ , while there was decreased IL-1 $\beta$ , IL-6, and IL-17 (Figure 5A). After *C. rodentium* infection, CXCL1, IL-6, IL-7, G-CSF, and IL-17 were increased in *Bves*<sup>-/-</sup> mice after adjusting for expression of these analytes at baseline (i.e. before infection, Figure 5B). Supplemental Table 1 summarizes the levels of all analytes after adjusting for baseline levels of expression. Of note, CXCL1 is chemotactic for polymorphonuclear cells; G-CSF stimulates survival, proliferation, and function of both neutrophils and neutrophil precursors; IL-6 is macrophage-derived and a pro-inflammatory marker; IL-7 is derived from dendritic cells and epithelial cells and stimulates proliferation of B cells, T cells, and NK cells. Finally, IL-17 is from Th17 cells and innate lymphoid cells, and most importantly is a well-established part of the pro-inflammatory response to *C. rodentium*.<sup>35</sup> Specific levels of protein expression are included in Supplemental Table 2 as well as all of the other chemokines and cytokines which were measured. Collectively, these observations suggest that *Bves*<sup>-/-</sup> colons are primed for a robust inflammatory response when exposed to *C. rodentium* and there is a significantly altered immune response post-infection.

### ***Bves*<sup>-/-</sup> mice exhibit upregulated IL-22 mediated mucosal host defenses after *C. rodentium***

We next examined expression of IL-22, which is known to be upregulated after *C. rodentium* infection. IL-22 deficient mice exhibit more severe colitis after *C. rodentium* and thus is thought to mediate the host's initial resistance.<sup>36</sup> Additionally, IL-22 is significantly upregulated in patients with Crohn's disease and Ulcerative Colitis.<sup>37</sup> *Il22* mRNA expression in whole colon after *C. rodentium* infection is increased in *Bves*<sup>-/-</sup> mice (Figure 6A). There is also significantly increased mRNA expression of IL-22 downstream targets, *Reg3b*, *Reg3g*, and *Nos2* in colonic epithelial isolates (Figure 6B). Additionally, immunofluorescence for neutrophils reveals increased neutrophilic infiltration in *Bves*<sup>-/-</sup> mice after *C. rodentium* infection (Figure 6C), consistent with increased CXCL1 expression observed in luminex profiling (Figure 5B). The increase in neutrophils, *Il22*, and its downstream targets suggests that host defense mechanisms could be upregulated in response to neutrophilic infiltration more directly and increased bacterial colonization and attachment to the epithelium more indirectly.

### ***Bves*<sup>-/-</sup> mice have worse colonic injury after DSS**

We next determined if BVES-dependent injury responses observed in infectious colitis modeling were generalizable to other methods of injury by treating WT and *Bves*<sup>-/-</sup> mice with DSS to induce a chemical colitis. Using colonoscopy to evaluate for colitis progression *Bves*<sup>-/-</sup> mice treated for 5 days with 4% DSS appeared to have more severe injury in comparison to WT control animals (Figure 7A). Histologic examination of colonic H&E staining and ulceration scores confirmed increased injury phenotypes in *Bves*<sup>-/-</sup> colons (Figure 7B). There was no difference in *Cxcl1* mRNA expression or neutrophilic infiltration in acute DSS trial (Supplementary Figure 5C–D). Notably, similar to what we observed in infectious injury modeling, *Bves*<sup>-/-</sup> mice did not exhibit a post-injury hyper-proliferative response. In fact, there was a decrease in proliferation when adjusting for number of proliferating cells before DSS (Figure 7C).

We next tested for repair process deficiencies by treating WT and *Bves*<sup>-/-</sup> mice with 4% DSS for 5 days and then allowed them to recover for 5 days (DSS replaced with water). While there was similar weight loss in both genotypes in both the acute and injury-repair DSS trials (Supplementary Figure 5A–B), histologic analysis determined that there was greater injury in *Bves*<sup>-/-</sup> mice (Figure 7D) and that proliferation was decreased in *Bves*<sup>-/-</sup> mice as compared with WT after the recovery period both before and after adjusting for pre-DSS proliferation (Figure 7E).

Finally, to model response to chronic epithelial injury, we treated WT and *Bves*<sup>-/-</sup> mice with 4% DSS and then allowed them to recover for 10 days, three times consecutively. After three cycles of 4% DSS, 100% of WT mice had survived, but only 62.5% of *Bves*<sup>-/-</sup> mice survived (Figure 7F). While there was not a significant difference in histologic ulceration score or proliferation in the surviving mice, this could potentially be due to loss of more moribund mice in the *Bves*<sup>-/-</sup> group, preventing histologic examination of the sicker mice. Thus, consistent with our results in the *C. rodentium* model of colitis, DSS treatment of *Bves*<sup>-/-</sup> mice yields greater histologic injury in acute and injury-repair modeling and increased mortality rates in a chronic model.

### **BVES limits EPEC infection in human cell lines**

To dissect the mechanism by which *Bves*<sup>-/-</sup> mice were more sensitive to colitogenic insult, we first studied the effect of EPEC, a *C. rodentium* human analog, on isolated epithelial cell cultures. We infected Caco2 cells, a polarized CRC-derived epithelial cell line, with EPEC as it has a similar mechanism of pathogenesis as *C. rodentium*, namely formation of attaching and effacing lesions and resultant epithelial damage. After knocking down *BVES* (Figure 8A) via an siRNA approach, 3 hour EPEC infection resulted in increased *CXCL8* expression (Figure 8B), which is known to be induced by EPEC infection and is an indicator of bacterial invasion.<sup>38</sup> There is also increased attachment of EPEC as evidenced by actin stalks (Figure 8C). These observations were replicated in HEK293T cells (Figure 8D). We observed a consistent phenotype with increased expression of *CXCL8* (Figure 8E) and increased attaching/effacing lesions per cell (Figure 8F) after knocking down *BVES*. Moreover, BVES transfected HEK293T cells (Supplemental Figure 6A) were then infected with EPEC for 3 hours. In these experiments, we observed decreased *CXCL8* expression (Supplemental Figure 6B) as well as a decreased number of attaching/effacing lesions by actin immunofluorescence in cells overexpressing BVES (Supplemental Figure 6C). These findings suggest that the pathogenicity of EPEC and hence *C. rodentium* can be limited by BVES expression.

### **BVES is underexpressed in moderate/severe ulcerative colitis**

Based on the above findings, we next examined clinical samples to determine if BVES levels were altered in ulcerative colitis. Clinical subjects were recruited from control and IBD subjects undergoing colonoscopy for evaluation of UC or non-UC indications at Vanderbilt University. Mucosal biopsies were obtained from control subjects and UC patients from areas of active disease, which was confirmed based on histological analysis.<sup>28</sup> RNA was extracted, qRT-PCR performed, and delta-delta Ct calculations between normal and disease samples indicated a 2.3-fold reduction of *BVES* mRNA levels in moderate/severe UC samples (Figure 9A). The patient characteristics, including age, gender, BMI, and current medications are shown in Figure 9B. These observations are consistent with our experimental data and suggest that BVES may also contribute to epithelial integrity in inflammatory bowel disease.

## **DISCUSSION**

In this study, we investigated the role of BVES in colonic homeostasis and in response to colon injury. At baseline, *Bves*<sup>-/-</sup> mice have a number of subtle colonic phenotypes which in aggregate we postulated might result in altered injury response when challenged. In every experimental model we tested, *Bves*<sup>-/-</sup> mice had exacerbated colitis and in the *C. rodentium* model there was increased bacterial translocation. EPEC infection of both Caco2 and HEK293T BVES knockdown cell lines increased the number of attaching/effacing lesions. Conversely, fewer attaching/effacing lesions occurred when *BVES* expression was increased in these lines, suggesting that BVES limited colonization. Lastly, *BVES* expression was reduced in ulcerative colitis samples, indicating relevance to human disease and a potential protective role for BVES in human disease.

We propose that colitis may be more severe in the absence of BVES due to compromised barrier function. Redistribution of phosphomyosin light chain (pMLC) to the luminal epithelial surface could potentially alter apical web contractile properties and thus increase permeability.<sup>39</sup> Interestingly, we had previously observed that when BVES function was impaired in LIM2405 cells pMLC expression increased.<sup>15</sup> Additionally, the downregulation of Claudin-7 and concurrent increase in *Claudin-2* could contribute to increased permeability and susceptibility to both infectious and chemical colitis.<sup>40,41</sup> Further, we observed increased expression of ZO-1<sup>42</sup> and JAM-A<sup>11</sup> which function to preserve epithelial barrier function. These increases could represent compensatory changes arising with BVES loss and potentially limit significant inflammation at baseline. However, these barrier defects may compromise the epithelium, priming for increased *C. rodentium* colonization and susceptibility to chemical colitis.

*C. rodentium* infection results in reproducible colonic inflammation and injury with fewer systemic effects in comparison to other models of infectious colitis.<sup>43,44</sup> Two pathognomonic features of *C. rodentium* infection include induction of actin cytoskeletal reorganization causing attaching/effacing lesions in the colonic epithelium as well as development of crypt hyperplasia, which is triggered by the loss of epithelial barrier integrity and the transit of bacteria into the lamina propria.<sup>45</sup> Knocking down BVES in both HEK293T and Caco2 cells increased the numbers of attaching/effacing lesions and increased *Cxcl8* expression, indicating more severe infection. One possible explanation for increased attaching/effacing lesions is that the loss of BVES and subsequent loss of cell-cell adhesion increases possible binding sites for EPEC and *C. rodentium*. *C. rodentium* infection also induces robust Th1 and Th17 immune responses.<sup>46,47</sup> After *C. rodentium* inoculation, *Bves*<sup>-/-</sup> mice demonstrated increased IL-17 protein expression and increased bacterial colonization and seeding of mesenteric lymph nodes, indicative of a more systemic infection. Moreover, increased *Il22*, *Reg3b*, *Reg3g*, and *Nos2* expression strongly suggests that host mucosal defenses were upregulated in response to more severe disease. As such, loss of BVES increases the susceptibility of colonic epithelial cells to attaching/effacing bacteria and upregulates immune responses.

It was unexpected that *Bves*<sup>-/-</sup> mice would have a hypoproliferative response after injury, especially in the setting of potentiated proliferation at baseline. While we don't have a unifying explanation for this phenotype at this time, one hypothesis is that injury resulted in a relative depletion of stem cells in *Bves*<sup>-/-</sup> mice resulting in reduced inflow into the transit amplifying cell compartment. This suggests that a role for BVES in stem cell preservation after injury. We speculate that increased stem cell activity at baseline places these populations at increased risk when injured.

Future experiments will further elucidate the mechanisms by which the stem cell compartment is compromised after chemical or infectious injury. We also plan to analyze the interaction between BVES and other tight junction proteins, particularly Claudin-7 and Claudin-2, which are dysregulated in *Bves*<sup>-/-</sup> mice. Additionally, our group has reported that BVES interacts with a subunit of PP2A,<sup>21</sup> which has previously been implicated in acetaldehyde-induced disruption of tight junctions in Caco2 monolayers.<sup>48</sup> We plan to

investigate the functional relevance of the BVES-PP2A interaction after injury *C. rodentium* and EPEC.

In conclusion, our findings demonstrate that BVES contributes to multiple aspects of intestinal homeostasis and colonic injury responses. Defects in other TJ constituents have compromised epithelial barrier function and lead to increased paracellular permeability,<sup>11,12,40,41</sup> similar to the phenotypes we observed with BVES loss. For example, loss of tight junction associated proteins JAM-A and NMIIA resulted in baseline inflammation and increased susceptibility to chemical colitis with DSS treatment.<sup>11,12</sup> Furthermore, TJ dysfunction is implicated in the pathogenesis of IBD. Prior studies have demonstrated that altered permeability due to tight junction disruption is associated with both increases in bacterial colonization and bacterial load, as well as hyperactive mucosal immune responses.<sup>49,50</sup> In these experiments, we identify a critical role for a tight junctional protein, BVES, in preserving colonic epithelial integrity under physiologic stress induced by both chemical and infectious agents. *Bves*<sup>-/-</sup> mice display increased permeability, have altered tight junction composition, have increased lymphoid aggregates, have worse colitis after DSS treatment, and have increased bacterial colonization and histologic injury after infection with *C. rodentium*. Finally, loss of *BVES* expression in Caco2 and HEK293T cells leads to increased attaching/effacing lesions after infection with EPEC, while these lesions are decreased with augmented *BVES* expression. These findings, combined with decreased expression of *BVES* in patients with UC, suggest that alterations in BVES may be relevant to understanding the mechanisms that underlie epithelial barrier dysfunction in colitis.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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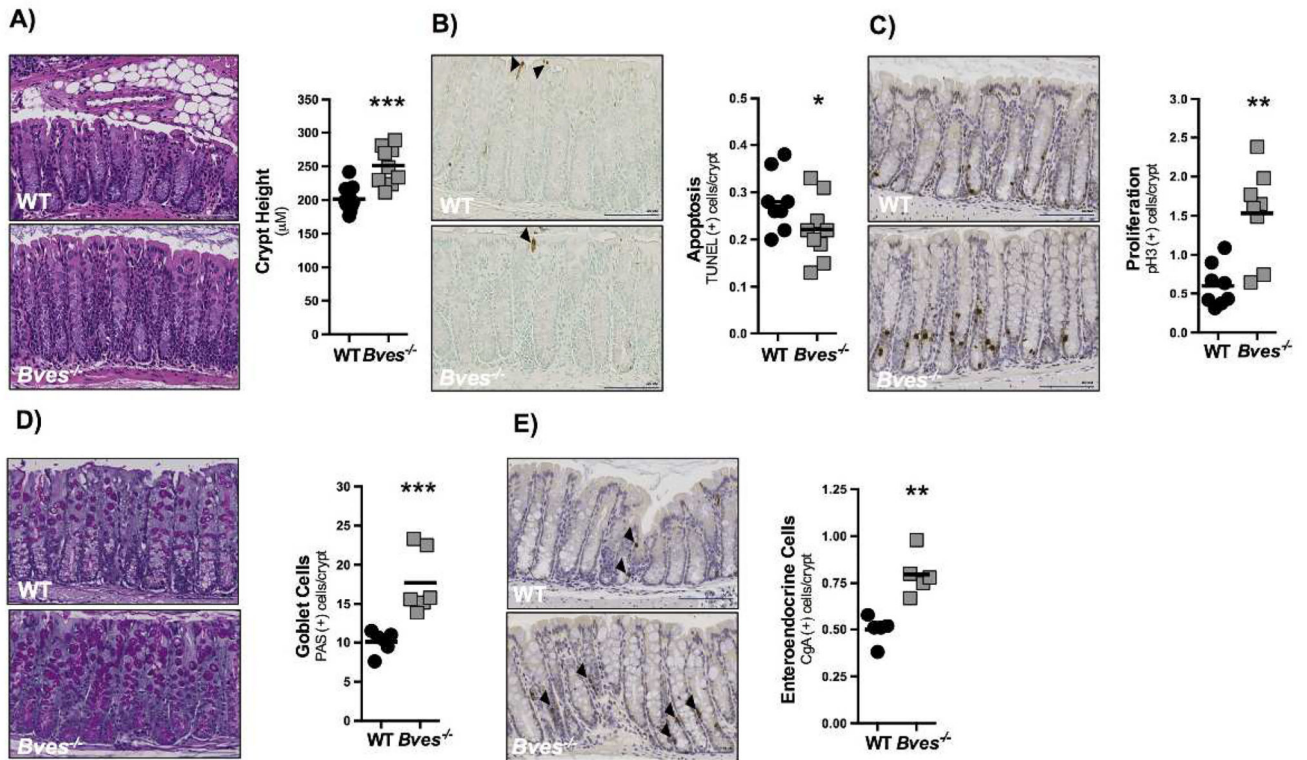
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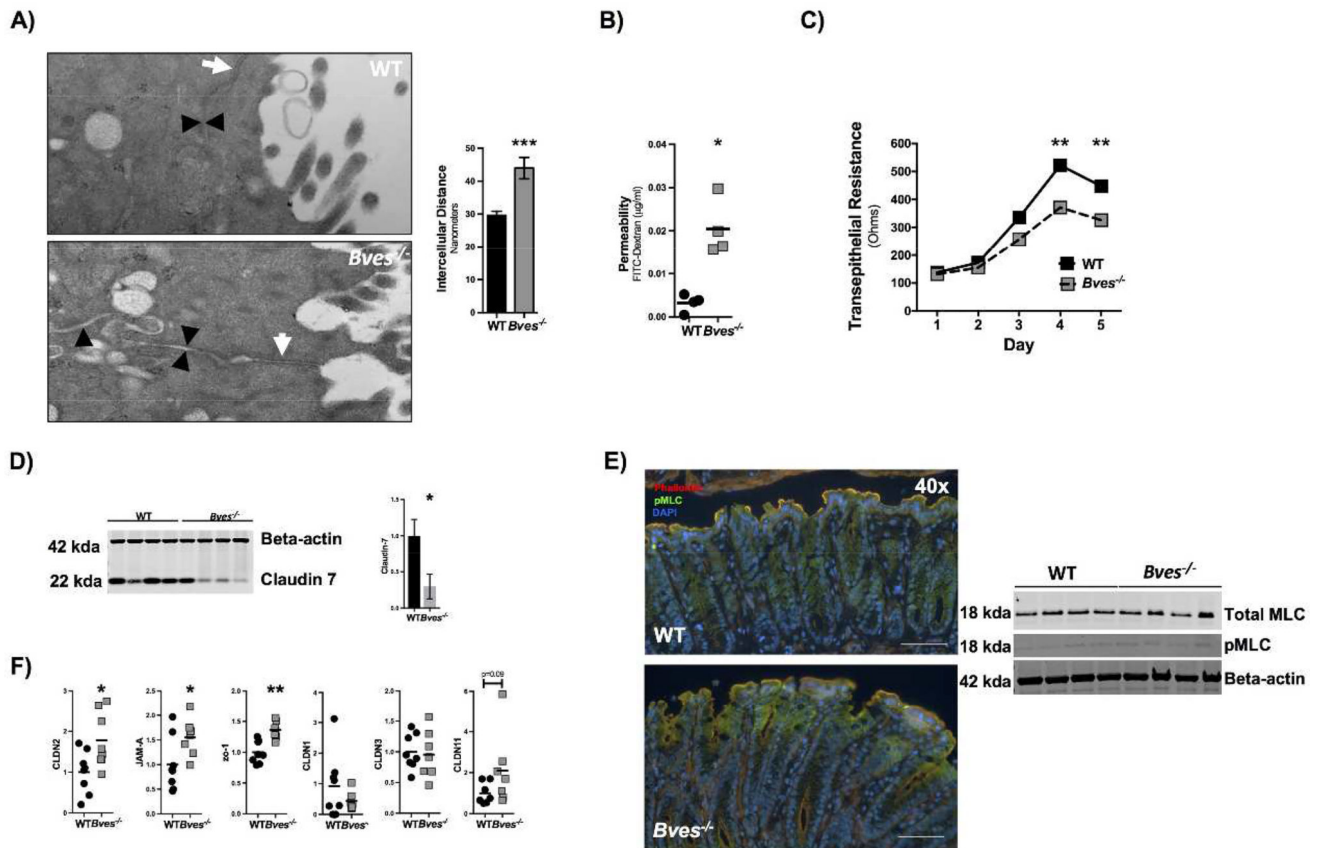
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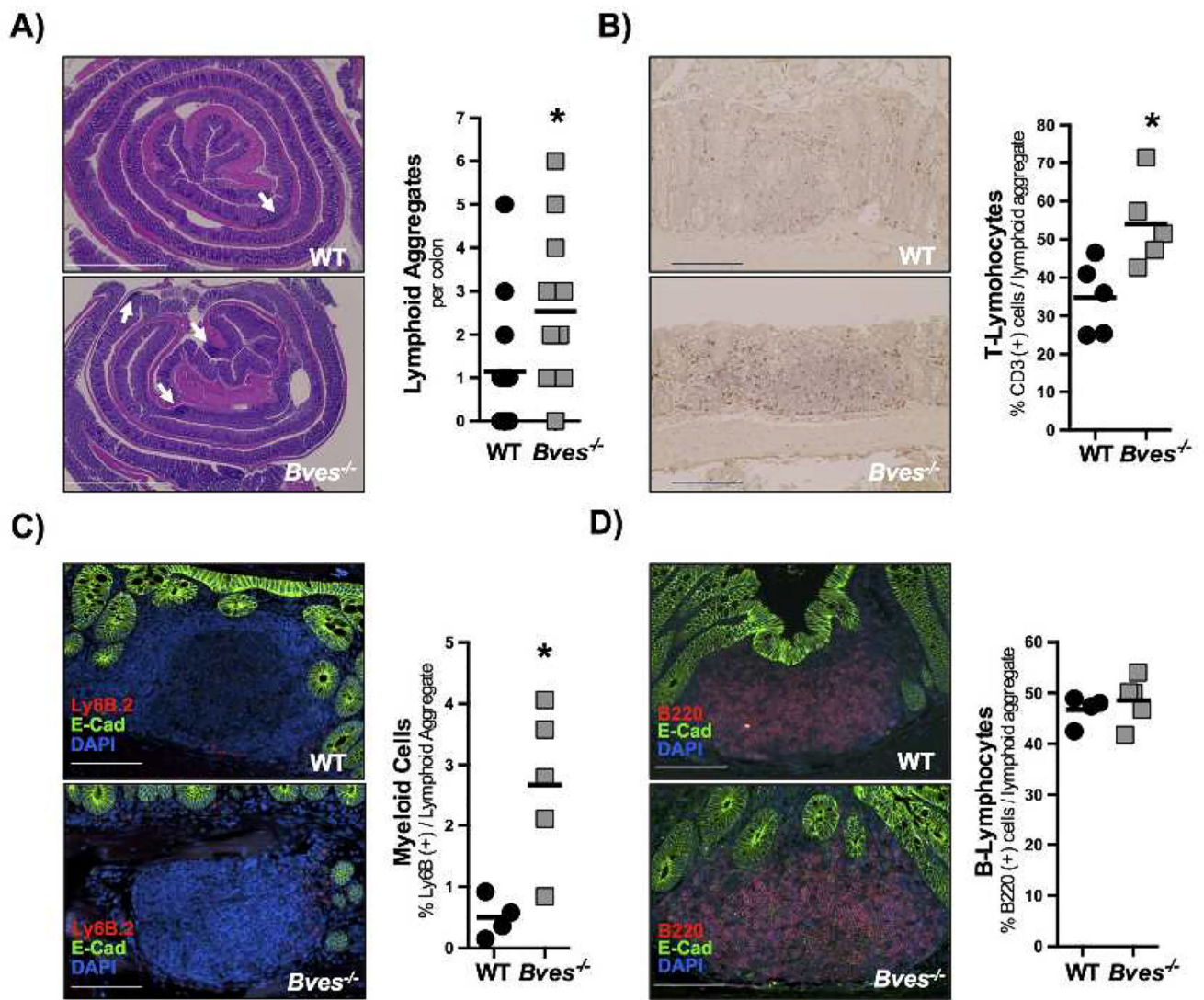


**Figure 1. BVES regulates colonic proliferation, lineage allocation, and crypt morphology**  
 Colons were isolated and Swiss-rolled. (A) Representative H&E staining of sections of WT and *Bves*<sup>-/-</sup> colons. Images (left) and quantification (right) of WT and *Bves*<sup>-/-</sup> crypt height (201.7  $\mu\text{m}$  vs. 251.2  $\mu\text{m}$ , \*\*\* $P$ <0.001,  $n$ =10). (B) Images (left) and quantification (right) of apoptotic cells per crypt unit (0.3 vs. 0.2 TUNEL<sup>+</sup> cells/crypt, \* $P$ =0.05,  $n$ =8). (C) Images (left) and quantification (right) of crypt proliferation (0.6 vs. 1.5 phospho-Histone H3<sup>+</sup> cells/crypt, \*\* $P$ <0.01,  $n$ =8). (D) Images (left) and quantification (right) of goblet cells/crypt (10.1 vs. 17.7 PAS<sup>+</sup> cells/crypt, \*\* $P$ <0.01,  $n$ =6). (E) Images (left) and quantification (right) of enteroendocrine cells/crypt-villus unit (0.5 vs. 0.8 CgA<sup>+</sup> cells/crypt, \*\* $P$ <0.01,  $n$ =5). All images were captured at 20 $\times$  magnification. Black arrows indicate positively-stained cells in (B) and (E). Scale bars are in the bottom right corner of each image and are 100  $\mu\text{m}$  in length.



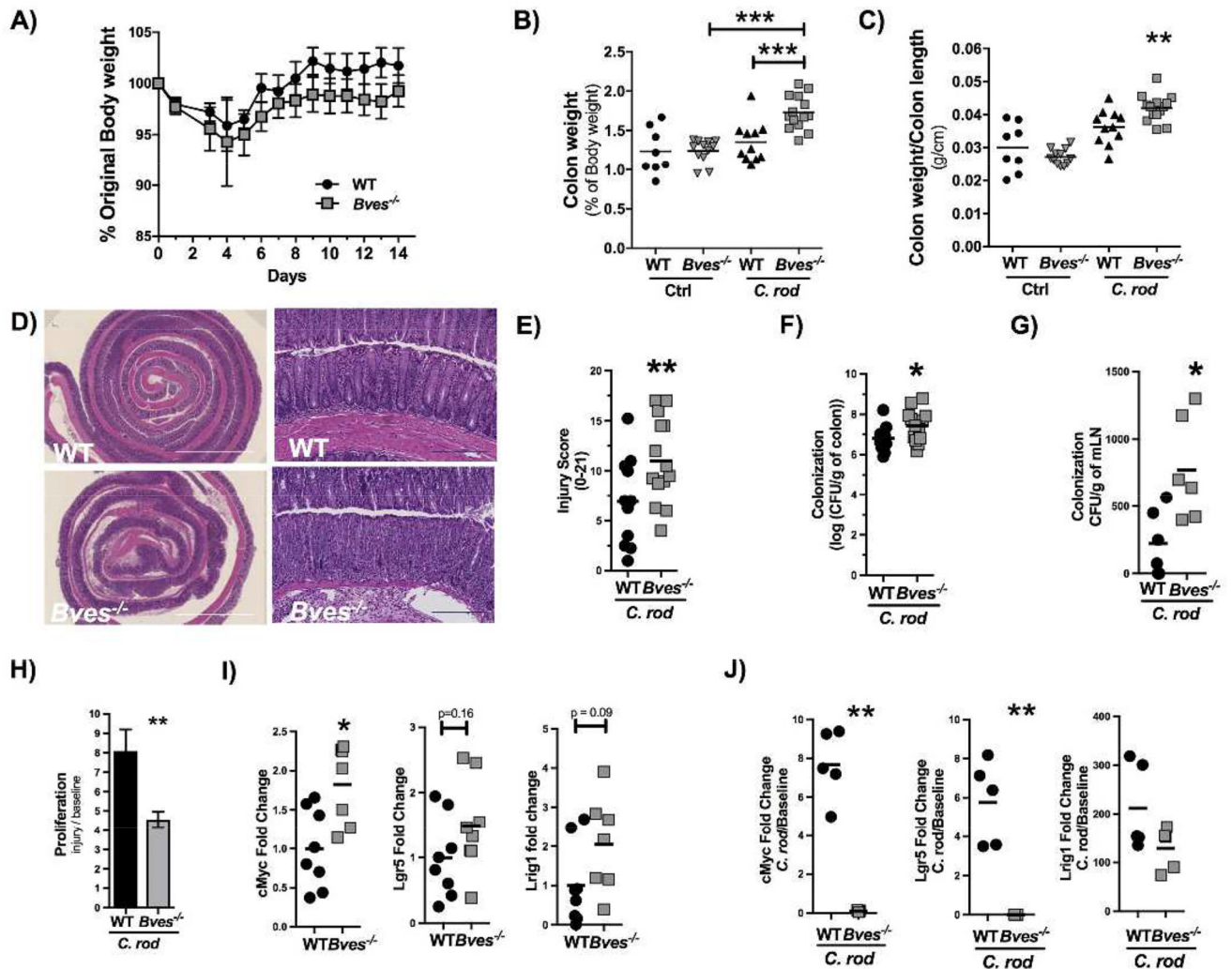
**Figure 2. *Bves*<sup>-/-</sup> colons display increased permeability and altered tight junction composition** (A) Intercellular spaces quantified by electron microscopy (*Bves*<sup>-/-</sup> vs. WT  $44.0 \pm 3.3$  nm vs.  $29.6 \pm 1.3$  nm, \*\*\* $P < 0.0001$ ,  $n = 100-105$  images from 3 mice). Images taken at 42,000x. White arrows label tight junctions and black arrowheads delineate intercellular spaces on the basolateral side. (B) Amount of penetrated FITC-dextran was quantified via Ussing chamber *ex vivo* permeability studies (*Bves*<sup>-/-</sup> vs. WT,  $0.020 \pm 0.003$  vs.  $0.003 \pm 0.001$  µg/mL, \* $P < 0.05$ ,  $n = 4$  in each group). (C) Colonoids were isolated and grown as monolayers on transwells, and transepithelial resistance (TER) was measured daily by Ohm meter. TER differences were quantified on each day, but only day 4 and day 5 exhibited significant differences: Day 4 (*Bves*<sup>-/-</sup> vs. WT,  $371.7 \pm 4.8$  vs.  $522.2 \pm 5.7$  ohms, \*\* $P < 0.01$ ,  $n = 6$ ) and day 5 ( $326.5 \pm 6.8$  vs.  $447.7 \pm 8.2$  ohms, \*\* $P < 0.01$ ,  $n = 6$ ). (D) Claudin 7 immunoblot from colonic crypts (left) and quantification (right) (*Bves*<sup>-/-</sup> vs. WT, 70% reduction, \* $P < 0.05$ ,  $n = 4$ ). (E) Immunofluorescence of the colonic epithelium shows redistribution of pMLC although pMLC and MLC immunoblot were nonsignificant ( $n = 4$ ). Scale bars are shown in the bottom right and are 100 µM. (F) Tight junction mRNA expression of *Bves*<sup>-/-</sup> and WT crypt isolates. *Claudin2* (*Bves*<sup>-/-</sup> vs. WT, 1.8 fold increase, \* $P < 0.05$ ,  $n = 7-8$ ); *JAM-A* (*Bves*<sup>-/-</sup> vs. WT, 1.6 fold increase, \* $P < 0.05$ ,  $n = 7-8$ ); *Zo-1* (*Bves*<sup>-/-</sup> vs. WT, 1.4 fold increase, \* $P < 0.01$ ,  $n = 7-8$ ); *Claudin1*, *Claudin3*, and *Claudin11* ( $P = ns$ ,  $n = 7-8$ ).





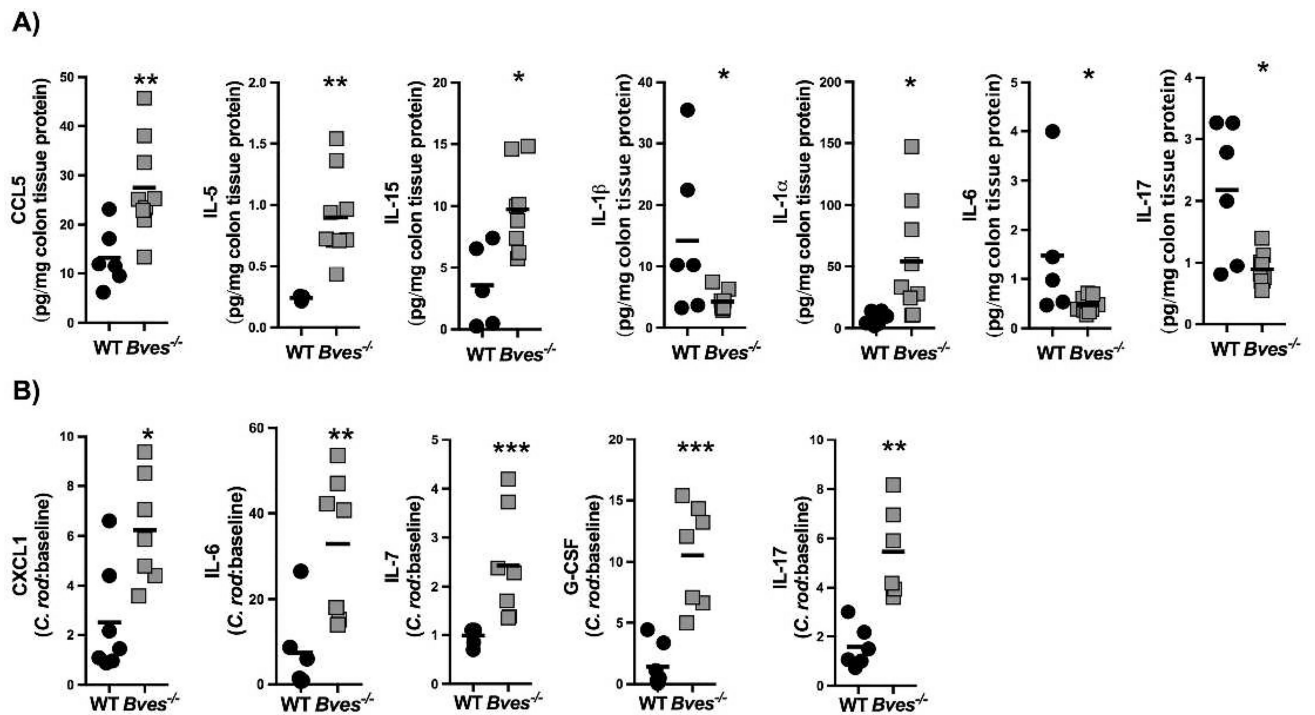
**Figure 3. *Bves*<sup>-/-</sup> colons have increased lymphoid aggregates**  
 Colons were isolated and swiss-rolled. For each analysis images are on left and quantification on right. (A) Total number of lymphoid aggregates per colon (*Bves*<sup>-/-</sup> vs. WT, 2.5 vs. 1.2, \**P* < 0.05, n=7-10). Scale bar in bottom left corner and is 500  $\mu$ m. (B) CD3 immunohistochemistry and CD3+ cells/lymphoid aggregate (*Bves*<sup>-/-</sup> vs. WT, 62.4 vs. 39.7, \**P* < 0.05, n=5) (C) Neutrophil immunofluorescent staining (Ly6B.2) and Ly6B.2+ cells/lymphoid aggregate (DAPI = blue, E-cadherin = green, and Ly6B.2 = red; *Bves*<sup>-/-</sup> vs. WT, 2.7% vs. 0.5%, \**P* < 0.05, n=4-5). (D) B cell immunofluorescence (B220) and B220+ cells/lymphoid aggregate (DAPI = blue, E-cadherin = green, and B220 = red; *Bves*<sup>-/-</sup> vs. WT, 48.6% vs. 46.7%, *P* = 0.56). For B-D, scale bars are in the bottom left corner and represent 100  $\mu$ m.





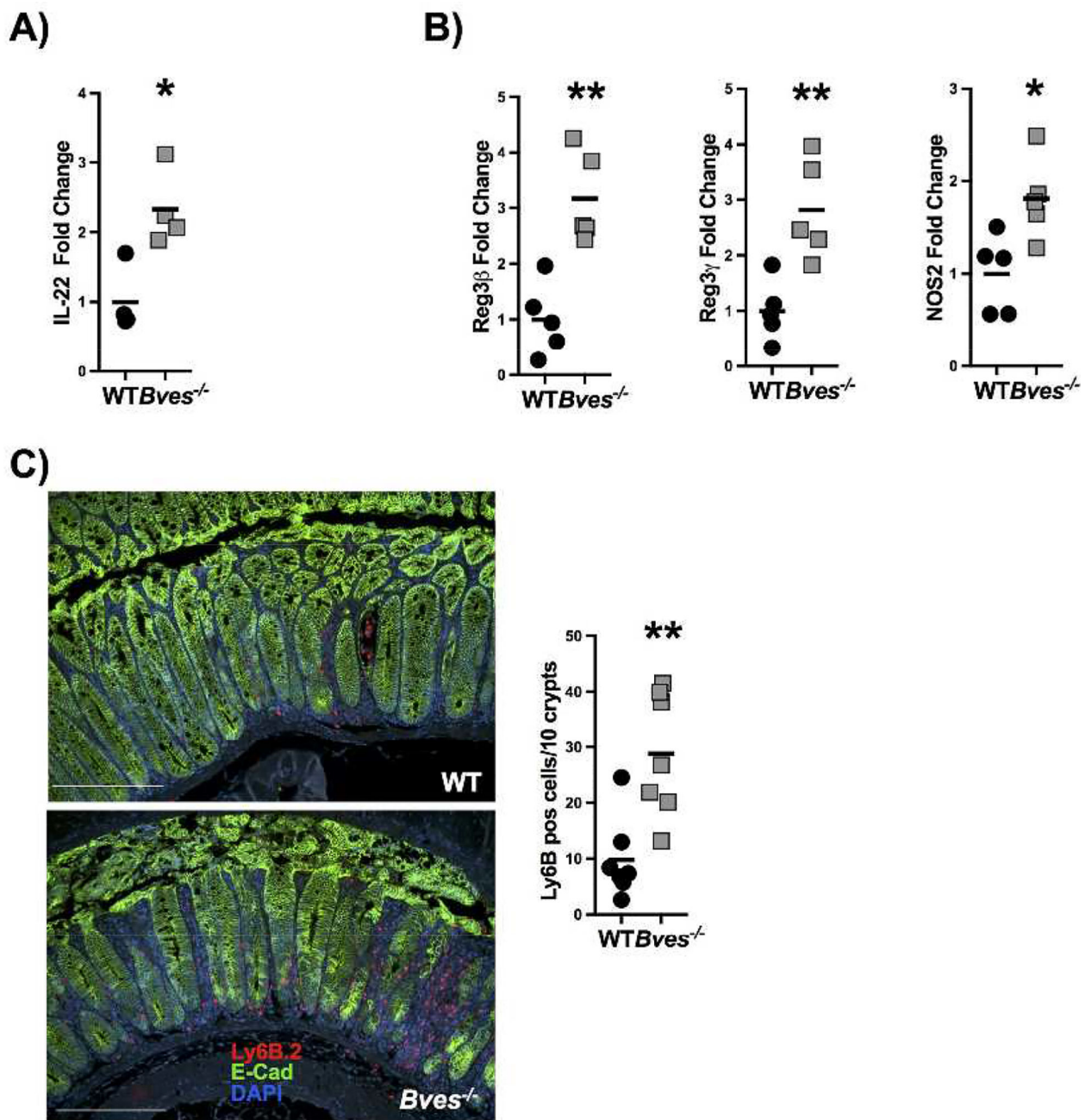
**Figure 4.** *Bves*<sup>-/-</sup> mice demonstrated augmented colitis after infection with *Citrobacter rodentium*. WT and *Bves*<sup>-/-</sup> mice were infected with *C. rodentium* and sacrificed 14 days after inoculation. (A) Body weight measured daily for 14 days after inoculation shown as a percentage of starting weight on each day (*Bves*<sup>-/-</sup> vs. WT, *P* = ns, *n* = 8–14). (B) Ratio of colon weight to body weight at the time of sacrifice (*Bves*<sup>-/-</sup> *C. rod* vs. WT *C. rod*,  $1.73 \pm 0.06$  vs.  $1.40 \pm 0.08$ , \*\*\**P* < 0.001, *n* = 8–14). (C) Ratio of colon weight to colon length at the time of sacrifice (*Bves*<sup>-/-</sup> vs. WT,  $0.04 \pm 0.001$  vs.  $0.03 \pm 0.002$ , \*\*\**P* < 0.001, *n* = 8–14). (D) Representative H&E stained sections of infected WT and *Bves*<sup>-/-</sup> mice. Scale bars are pictured in the lower left hand corner. They represent 500  $\mu$ M for 5 $\times$  images on left, while they represent 100  $\mu$ M for the 20 $\times$  images on the right. (E) Injury scores (*Bves*<sup>-/-</sup> vs. WT,  $6.9 \pm 1.3$  vs.  $11.0 \pm 1.1$ , \*\**P* < 0.01, *n* = 11–14). Histological injury score using a multi-point scale as described in the Methods section. (F) Bacterial colonization of colon with *C. rodentium* quantified as log (CFU/g colon) (*Bves*<sup>-/-</sup> vs. WT,  $7.4 \pm 0.4$  vs.  $6.8 \pm 0.2$ , \**P* < 0.05, *n* = 8–13). (G) Bacterial colonization of mesenteric lymph nodes (mLNs) calculated as CFU/g mLN (*Bves*<sup>-/-</sup> vs. WT,  $224 \pm 99$  vs.  $773 \pm 156$  CFU/g, \**P* < 0.05, *n* = 6). (H) Colonocyte proliferation was calculated as a ratio of the number of pH3 positive stained

cells after treatment with *C. rodentium* divided by number of pH3 positive stained cells at baseline (*Bves*<sup>-/-</sup> Injury/Baseline vs. WT Injury/Baseline,  $4.5 \pm 0.4$  vs.  $8.1 \pm 3.7$ , \*\* $P < 0.01$ ,  $n = 11-13$ ). (I) mRNA expression of stem cell markers from colon epithelial isolates at baseline quantified by qPCR: *cMyc* (*Bves*<sup>-/-</sup> vs. WT,  $1.8 \pm 0.2$  vs.  $1.0 \pm 0.2$  fold change, \* $P < 0.05$ ,  $n = 7-8$ ), *Lgr5* (*Bves*<sup>-/-</sup> vs. WT,  $1.5 \pm 0.3$  vs.  $1.0 \pm 0.2$   $P = 0.16$ ,  $n = 8$ ) and *Lrig1* (*Bves*<sup>-/-</sup> vs. WT,  $2.1 \pm 0.5$  vs.  $1.0 \pm 0.4$ ,  $P = 0.09$ ,  $n = 7-8$ ). (J) After *C. rodentium* infection, expression of stem cell markers relative to baseline was quantified by qPCR. *cMyc* (*Bves*<sup>-/-</sup> vs. WT,  $0.1 \pm 0.02$  vs.  $7.7 \pm 0.8$  fold change relative to baseline, \*\* $P < 0.01$ ), *Lgr5* (*Bves*<sup>-/-</sup> vs. WT,  $0.004 \pm 0.0005$  vs.  $5.8 \pm 0.9$  fold change relative to baseline, \*\* $P < 0.01$ ), and *Lrig1* (*Bves*<sup>-/-</sup> vs. WT,  $75 \pm 20$  vs.  $212 \pm 40$  fold change relative to baseline,  $P = 0.42$ ).



**Figure 5. *Bves*<sup>-/-</sup> colons display dysregulated chemokine and cytokine expression by Luminex multi-analyte profiling**

(A) Chemokines and cytokines which were significantly different between WT and *Bves*<sup>-/-</sup> colon at baseline: CCL5 (\*\* $P$ <0.01,  $n$ =7–8), IL-1 $\alpha$  (\* $P$ <0.05,  $n$ =6–8), IL-5 (\*\* $P$ <0.01,  $n$ =3–7), IL-1 $\beta$  (\* $P$ <0.05), IL-15 (\* $P$ <0.05,  $n$ =5–6), IL-6 (\* $P$ <0.05,  $n$ =5–8), and IL-17 (\* $P$ <0.05,  $n$ =6–8). (B) Chemokines and cytokines which are significantly different between WT and *Bves*<sup>-/-</sup> colons after *C. rodentium*: CXCL1 (\* $P$ <0.05,  $n$ =7), IL-6 (\*\* $P$ <0.01,  $n$ =6–7), IL-7 (\*\*\* $P$ <0.001,  $n$ =6–7), G-CSF (\*\*\* $P$ <0.001,  $n$ =7), and IL-17 (\*\* $P$ <0.01,  $n$ =6).



**Figure 6.** *Bves*<sup>-/-</sup> mice exhibited upregulated IL-22 mediated mucosal host defenses after *C. rodentium* (A) *Il22* mRNA expression in whole colon after *C. rodentium* infection was quantified by qPCR (*Bves*<sup>-/-</sup> vs. WT, 2.3 ± 0.3 vs. 1.0 ± 0.2 fold change, \**P*<0.05, n=4). (B) IL-22 downstream targets are also quantified by qPCR in colonic epithelial isolates: *Reg3b* (*Bves*<sup>-/-</sup> vs. WT, 3.2 ± 0.4 vs. 1.0 ± 0.3 fold change, \*\**P*<0.01, n=5), *Reg3g* (*Bves*<sup>-/-</sup> vs. WT, 2.8 ± 0.4 vs. 1.0 ± 0.2 fold change, \*\**P*<0.01, n=5), and *Nos2* (*Bves*<sup>-/-</sup> vs. WT, 1.8 ± 0.2 vs. 1.0 ± 0.2 fold change, \**P*<0.05, n=5). (C) Neutrophils were quantified by immunofluorescence for Ly6B.2 (*Bves*<sup>-/-</sup> vs. WT, 28.8 ± 4.2 vs. 9.8 ± 2.7 neutrophils per 10 crypts, \*\**P*<0.01,

n=7). Images shown are at 20x. Scale bars are pictured in the lower right-hand corner. They represent 100  $\mu\text{m}$ .

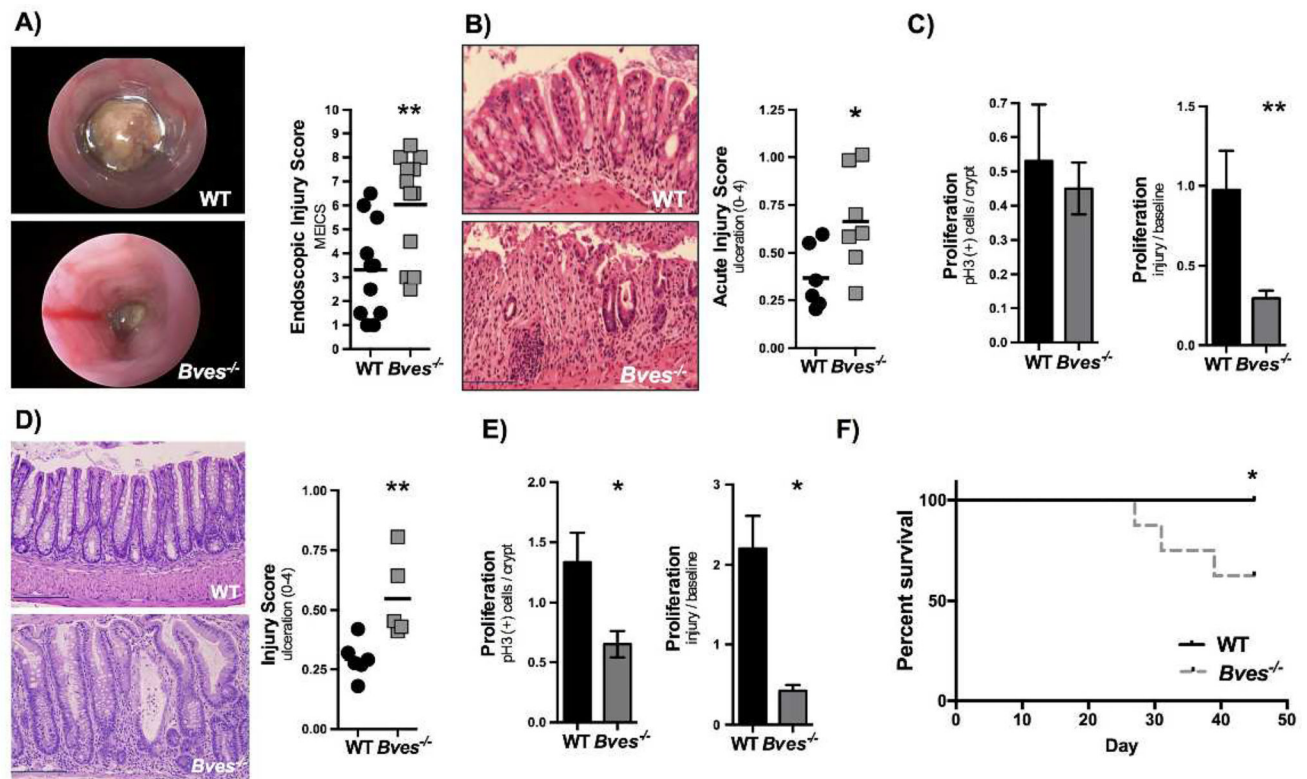
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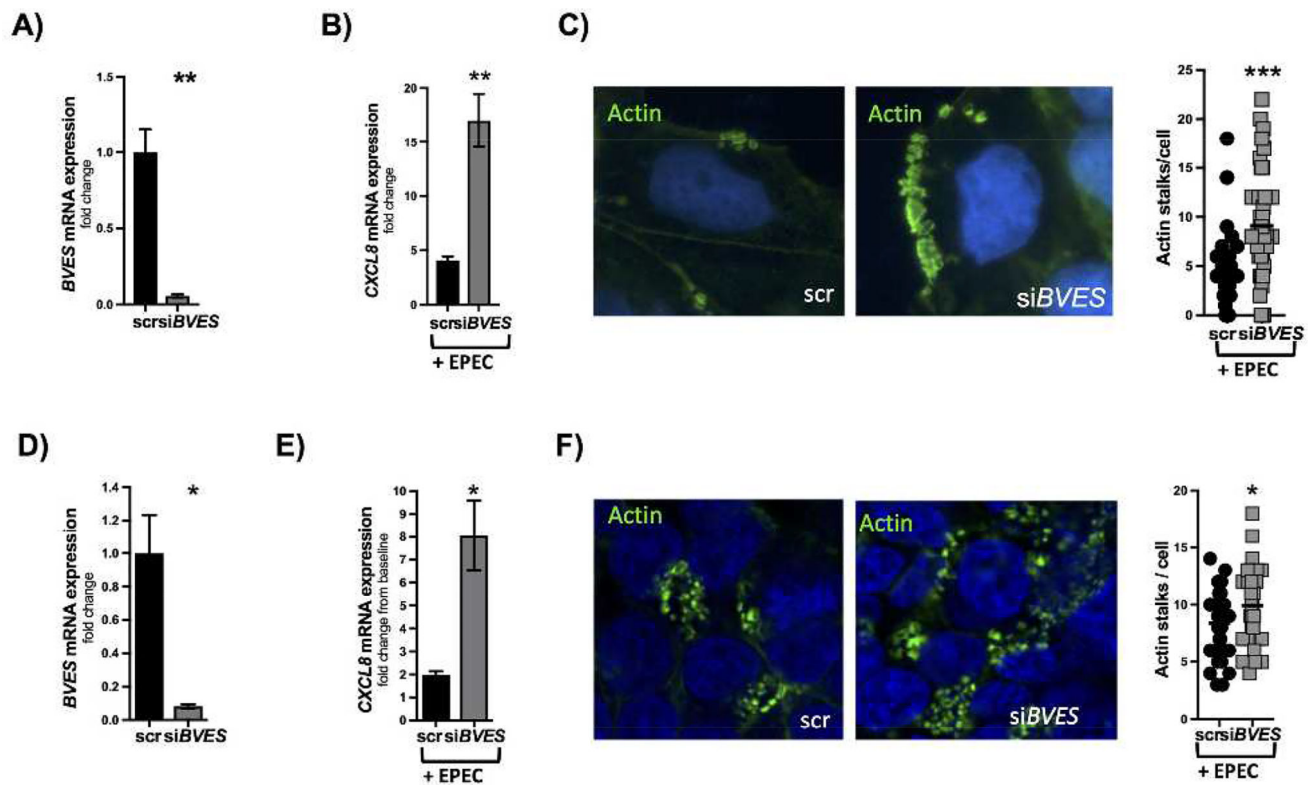




**Figure 7. *Bves*<sup>-/-</sup> mice demonstrated more severe colitis after DSS treatment**

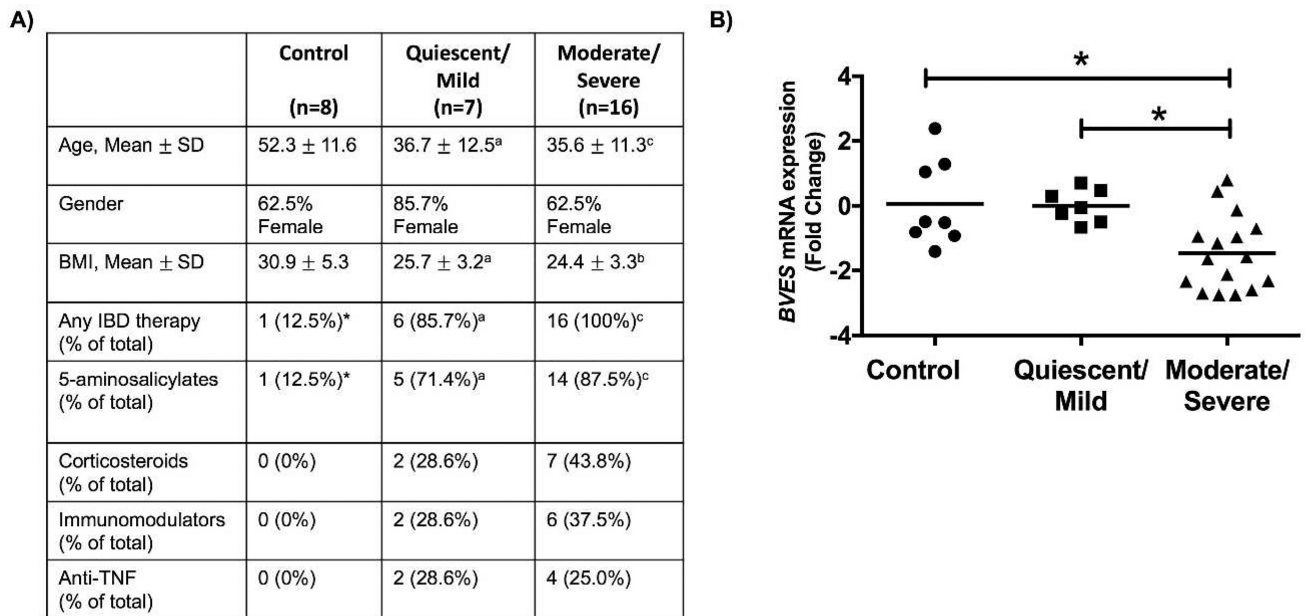
(A) WT (n=10) and *Bves*<sup>-/-</sup> mice (n=13) were treated with 4% DSS for 5 days and then sacrificed. Colonoscopy was performed prior to sacrifice. Colonoscopy videos were scored using MEICS (*Bves*<sup>-/-</sup> vs. WT,  $6.0 \pm 0.6$  vs.  $3.3 \pm 0.6$ , \*\* $P < 0.01$ , n=10–13). (B) Acute injury scores quantified after sacrifice (*Bves*<sup>-/-</sup> vs. WT,  $0.7 \pm 0.1$  vs.  $0.4 \pm 0.07$ , \* $P < 0.05$ , n=6–7). Scale bars are pictured in the lower left hand corner. They represent 100  $\mu$ M. (C) Left: Proliferation was defined by pH3 positive stained cells/crypt in the WT mice as compared with *Bves*<sup>-/-</sup> mice (*Bves*<sup>-/-</sup> vs. WT,  $0.45 \pm 0.08$  vs.  $0.53 \pm 0.2$ ,  $P = \text{ns}$ , n=7). Right: Proliferation was calculated as a ratio of the number of pH3 positive stained cells after treatment with DSS divided by number of pH3 positive stained cells at baseline (*Bves*<sup>-/-</sup> Injury/Baseline vs. WT Injury/Baseline,  $0.3 \pm 0.05$  vs.  $1.0 \pm 0.2$ , \*\* $P < 0.01$ , n=6–7). (D) Injury-repair model. Mice were treated with 4% DSS for 5 days and then allowed to recover for 5 days. The mice were then sacrificed. There was significantly greater histologic injury in the *Bves*<sup>-/-</sup> mice ( $0.6 \pm 0.08$  vs.  $0.3 \pm 0.03$ , \*\* $P < 0.01$ , n=4–5). Scale bars are pictured in the lower left hand corner. They represent 100  $\mu$ M. (E) Left: Increased proliferation defined by pH3 positive stained cells/crypt in the WT mice as compared with *Bves*<sup>-/-</sup> mice (*Bves*<sup>-/-</sup> vs. WT,  $0.65 \pm 0.1$  vs.  $1.3 \pm 0.2$ , \* $P < 0.05$ , n=4–5). Right: There was also increased proliferation when compared with mice not treated with DSS (i.e. when compared with baseline proliferation) (*Bves*<sup>-/-</sup> Injury/Baseline vs. WT Injury/Baseline,  $0.4 \pm 0.1$  vs.  $2.2 \pm 0.4$ , \* $P < 0.05$ , n=4–5). (F) Survival curves from mice treated with 3 cycles of 4% DSS for 5 days and 10 days of recovery (*Bves*<sup>-/-</sup> vs. WT, 62.5% vs. 100%). Survival curves were compared using Log-rank (Mantel-Cox) test. \* $P < 0.05$ , n=8–9.





**Figure 8. BVES influences pathogenicity of EPEC in Caco2 and HEK cells**

(A) *BVES* expression after siRNA knockdown in Caco2 cells (si*BVES* vs. scr,  $0.06 \pm 0.01$  vs.  $1.0 \pm 0.16$ ,  $**P < 0.01$ ). (B) *CXCL8* mRNA expression quantified after 3 hours EPEC infection in Caco2 cells (si*BVES* vs. scr,  $17 \pm 4.5$  fold vs.  $4.1 \pm 0.4$  fold,  $**P < 0.01$ , 2 experiments with 3 replicates each time). (C) Number of actin stalks per Caco2 cell after 3 hours EPEC infection (si*BVES* vs. scr,  $9.1 \pm 0.9$  vs.  $4.0 \pm 0.6$ ,  $***P < 0.001$ , 2 experiments with 3 replicates each time). Green dots represent individual actin stalks. Images are taken at 80x. (D) *BVES* knockdown by an siRNA approach in HEK293T cells (scr vs. si*BVES*,  $1.0 \pm 0.2$  vs.  $0.08 \pm 0.01$ ,  $*P < 0.05$ ). (E) *CXCL8* expression mRNA expression quantified after 3 hours EPEC infection in HEK293T cells (scr vs. si*BVES*,  $2.0 \pm 0.2$  fold vs.  $8.1 \pm 1.5$  fold,  $*P < 0.05$ , 2 experiments with 3 replicates each time). (F) Number of actin stalks per HEK293T cell quantified after 3 hours EPEC infection (scr vs. si*BVES*,  $8.4 \pm 0.5$  vs.  $9.9 \pm 0.5$ ,  $*P < 0.05$ ).



**Figure 9. *BVES* is underexpressed in moderate to severe ulcerative colitis**

*BVES* mRNA levels were determined using qRT-PCR on UC biopsy specimens. (A) UC status determined by histologic status. <sup>a</sup> $P < 0.05$ , <sup>b</sup> $P < 0.01$ , <sup>c</sup> $P < 0.001$  vs. control. \*This patient was started on therapy for presumed UC but with normal endoscopic evaluation was felt to have irritable bowel syndrome. Age and body mass index statistics were assessed by the Kruskal-Wallis test followed by the Mann-Whitney test. (B) *BVES* mRNA levels expressed as negative ddCt in normal, quiescent/mild UC, and moderate/severe UC. *BVES* expression quantified between normal and moderate/severe UC (\* $P < 0.05$ ,  $n = 7-8$ , Kruskal-Wallis) and between quiescent/mild and moderate/severe UC (\* $P < 0.05$ ,  $n = 7-16$ , Kruskal-Wallis).