Differential Effects of β-catenin and NF-κB Interplay in the Regulation of Cell Proliferation, Inflammation and Tumorigenesis in Response to Bacterial Infection

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Abstract

Both β-catenin and NF-κB have been implicated in our laboratory as candidate factors in driving proliferation in an in vivo model of Citrobacter rodentium (CR)-induced colonic crypt hyper-proliferation and hyperplasia. Herein, we test the hypothesis that β-catenin and not necessarily NF-κB regulates colonic crypt hyperplasia or tumorigenesis in response to CR infection. When C57Bl/6 wild type (WT) mice were infected with CR, sequential increases in proliferation at days 9 and 12 plateaued off at day 19 and paralleled increases in NF-κB signaling. In Tlr4−/− (KO) mice, a sequential but sustained proliferation which tapered off only marginally at day 19, was associated with TLR4-dependent and independent increases in NF-κB signaling. Similarly, increases in either activated or total β-catenin in the colonic crypts of WT mice as early as day 3 post-infection coincided with cyclinD1 and c-myc expression and associated crypt hyperplasia. In KO mice, a delayed kinetics associated predominantly with increases in non-phosphorylated (active) β-catenin coincided with increases in cyclinD1, c-myc and crypt hyperplasia. Interestingly, PKCζ-catalyzed Ser-9 phosphorylation and inactivation of GSK-3β and not loss of wild type APC protein accounted for β-catenin accumulation and nuclear translocation in either strain. In vitro studies with Wnt2b and Wnt5a further validated the interplay between the Wnt/β-catenin and NF-κB pathways, respectively. When WT or KO mice were treated with nanoparticle-encapsulated siRNA to β-catenin (si- β-Cat), almost complete loss of nuclear β-catenin coincided with concomitant decreases in CD44 and crypt hyperplasia without defects in NF-κB signaling. si-β-Cat treatment to ApcMin/+ mice attenuated CR-induced increases in β-catenin and CD44 that halted the growth of mutated crypts without affecting NF-κB signaling. The predominant β-catenin-induced crypt proliferation was further validated in a Castaneus strain (B6.CAST.11M) that exhibited significant crypt hyperplasia despite an attenuated NF-κB signaling. Thus, β-catenin and not necessarily NF-κB regulates crypt hyperplasia in response to bacterial infection.

Introduction

Both β-catenin and NF-κB proteins are important regulators of gene expression and cellular proliferation. β-catenin is a ubiquitously expressed protein that acts as an adhesion molecule by mediating the association of E-cadherin with the cytoskeleton [1,2] and is also a critical downstream component of the Wnt pathway [3–5]. The Wnt family of secretory glycoproteins plays an important role in embryonic development, in the induction of cell polarity, and in the determination of cell fate. Most mammalian genomes including human, harbor 19 Wnt genes, and is also a critical downstream component of the Wnt pathway mediating the association of E-cadherin with the cytoskeleton [1,2]. Wnt signaling can be broadly categorized as canonical if they require signaling via β-catenin while non-canonical Wnt signaling pathways are β-catenin independent. Interestingly, multiple studies have demonstrated a regulatory cross-talk between canonical and non-canonical pathways wherein, Wnt5a has been shown to induce a non-canonical pathway that inhibits canonical Wnt signaling. This observation has been made both in vivo and in vitro, including hematopoietic stem cells [8,9]. The outcome however, depends on the receptor context [10]. Wnt5a inhibits β-catenin-dependent activity when its signal is mediated by the orphan tyrosine kinase receptor Ror2 [11]. However, it activates β-catenin-dependent signaling when it signals through the co-receptors Fzd4 and LRP5.

NF-κB comprises a family of transcription factors which are critical in activating the expression of genes involved in the immune and inflammatory response and in the regulation of cellular apoptosis [12,13]. Typically, NF-κB is activated by inflammatory mediators such as TNF-α and IL-1β, via activation of IκB kinases (IKKs), which carry out the phosphorylation-dependent degradation of IκB inhibitors upon inflammatory...
stimuli [14–18]. Constitutively active NF-κB plays an important role in cancer progression and metastasis and has been reported to physically interact with β-catenin, resulting in a reduction in NF-κB nucleus translocation, DNA binding and target gene expression. Recently, GSK-3β is shown to be required for NF-κB activation [19]. As β-catenin is a major substrate of GSK-3β, it raises an interesting possibility that β-catenin may mediate the cross-regulation between the two pathways. Interestingly, β-catenin-independent Wnt signaling is also involved in various biological functions, such as vertebral development, cell motility and adhesion, and cancer invasiveness [20–23]. Indeed, Wnt5a has been recently implicated in certain inflammatory diseases [24,25]. Wnt5a expression is dependent on NF-κB signaling via Toll-like receptor (TLR) in human antigen-presenting cells [25,26]. Furthermore, IL-6 family members activate the gp130-JAK-signal transducer and activator of transcription 3 (STAT3) signaling cascade to up-regulate Wnt5a transcription in chronic persistent inflammation and rheumatoid arthritis [27]. However, little is known about the expression and modulation of non-canonical Wnt signaling in the intestine in response to bacterial infection.

It has been shown previously that development of necrotizing enterocolitis (NEC), a leading cause of death from gastrointestinal disease in preterm infants, requires activation of TLR4 on enterocytes and that TLR4 activation by bacterial endotoxin LPS (lipopolysaccharide) inhibits enterocyte proliferation and leads to disease progression due to impaired β-catenin signaling after activation of the Akt-GSK3β signaling pathway [28–30]. These effects were however, restricted to the small intestine of newborn mice, suggesting relevance in the development of NEC. Whether TLR4-mediated inhibition of β-catenin signaling in response to infection by Gram negative bacteria is a global phenomenon is not known.

Citrobacter rodentium (CR) naturally infects mice using a mechanism similar to those employed by attaching and effacing (A/E) bacterial pathogens EPEC and EHEC [31,32]. CR is an A/E pathogen which causes increased proliferation in the distal colon of adult outbred mice without associated injury or significant histological inflammation [33]. In genetically susceptible strains, clinical signs such as retarded growth, diarrhea, dehydration, coat ruffling, hunched posture and high mortality have been reported [33]. Utilizing the CR-infection model, we showed for the first time that colonic crypt hyperplasia was associated with NF-κB activation [34] and alterations in Casein Kinase-Ie that influenced β-catenin signaling [35,36]. Ongoing studies from our laboratory have further demonstrated that functional cross-talk between Wnt/β-catenin and Notch [37] and Notch and NF-κB [38] pathways regulate crypt hyperplasia and/or tumorigenesis in response to CR infection in outbred mice while inflammation and/or colitis in the inbred mice, driven by the expression of distinct cytokines/chemokines, is regulated by activation of the MEK/ERK/NF-κB pathways [39]. It was shown previously that TLR4 signaling contributes to inflammation induced by CR [40]. Based on the recent findings that TLR4 antagonizes β-catenin-induced cell proliferation in the small intestine but not in the colon, we hypothesized that β-catenin and not necessarily NF-κB will dictate the colonic crypt hyperplastic response following CR infection in mice deficient for Tlr4. This hypothesis was tested in the current study.

Results

Infection of C57Bl/6 wild type mice with Citrobacter rodentium (CR) elicited a predictable response in the distal colon: gross thickening accompanied by hyperplasia and significant increase in crypt length between days 7 to 12 post-infection. The crypt length however, plateaued between days 12 to 19 (Fig. 1A). To determine if changes in epithelial cell proliferation contributed towards differences in crypt lengths, we next stained colonic sections for Ki-67 as a marker for proliferation. Representative sections from the distal colons of uninfected normal (N) mice and from days 3 to 19 post-infected mice are shown in Fig. 1B. In normally proliferating crypts, only cells at the base exhibited nuclear staining (Fig. 1B). Between days 3–7 post-infection, a gradual increase in Ki-67 staining was recorded which peaked by day 9 before tapering off between days 12–19 (Fig. 1B).

Infection of Tlr4−/− mice with CR elicited a much more profound response both in terms of gross morphology and crypt epithelial cell proliferation (Fig. 1C, D). Interestingly, neither the crypt length nor the cell proliferation in response to CR infection decreased between days 7–19. On the contrary, Ki-67 staining even at day 19 (Fig. 1D) was significantly higher than the wild type counterpart (see Fig. 1B) suggesting a more aggressive response to infection in the absence of functional TLR4.

We have recently shown that NF-κB activation in response to CR infection involves signaling via TLR4 [41]. To determine if CR infection affected changes in TLR4 levels and to definitively characterize TLR4’s role in NF-κB activation during TMCH, we began by examining sequential changes in TLR4 in the colonic crypts of C57Bl/6J mice. As shown in Fig. 2A, TLR4 levels started to increase by day 3 and peaked between days 5–12 compared to uninfected control before declining at day 19. During measurement of NF-κB activity in the crypt nuclear extracts, a sequential increase was recorded between days 3–12 before activity declined at day 19 (Fig. 2B). We next investigated the phosphorylation status of p65-NF-κB to determine whether post-translational modification of this subunit contributed towards sustained activation of NF-κB following CR infection. When we utilized antibodies specific for Serines-276 (p65276) and Serines-536 (p65536) phosphorylation of p65 subunit, relative levels of both cellular and nuclear p65276 and p65536 along with total p65 increased significantly between days 3–12 compared with uninfected control while relative levels of both moieties declined at day 19 (Fig. 2C) and correlated with NF-κB activity and expression of its downstream target CXCL1-1/KC at these time points (Fig. 2D). The bar graph in Fig. 2D represents the ratio of CXCL1-1 normalized to actin from three independent experiments. These studies corroborate well with other murine models of IBD [42] suggesting that increases in TLR4 may be critical in regulating NF-κB activity during CR-induced crypt hyperplasia.

When Tlr4−/− mice were infected with CR for 3–19 days, there was no detectable TLR4 in the crypts of these mice as expected (Fig. 2E). During measurement of NF-κB activity in the crypt nuclear extracts, we did not observe any change at day 3 post-infection suggesting that absence of TLR4 apparently affected early activation of NF-κB in response to CR infection. At days 5 and 12 however, a bi-phasic activation of NF-κB was recorded followed by a declining trend at day 19 (Fig. 2F). Western blots with antibodies to p65276 and p65536 at days 5 and 12 dramatically correlated with NF-κB activation kinetics while changes in relative cellular and nuclear levels at other time points were less significant (Fig. 2G). The changes in NF-κB activity at days 5, 12 and 19 correlated well with CXCL1-1/KC kinetics (Fig. 2H). The bar graph in Fig. 2H represents the ratio of CXCL1-1 normalized to actin from three independent experiments. These studies clearly suggest that while early NF-κB activation in response to CR infection of Tlr4−/−
mice requires signaling via TLR4, later time points exhibit a TLR4-independent mechanism of NF-κB activation in response to CR infection. Given that TLR4 plays an important role in protective immunity wherein, it limits bacterial dissemination to mesenteric lymph nodes and peripheral organs such as liver and spleen, we next measured bacterial counts in the distal colons of uninfected and CR-infected *Tlr4*2/2 mice. As depicted in Fig. 2I, we did not measure any appreciable increase in bacterial counts at day 9 post-infection. At day 12, however, significant increase in bacterial counts compared to control, were recorded followed by a decline at day 19 (Fig. 2I). Whether the increase in bacterial counts at day 12 also represents bacterial translocation to subepithelial compartments is currently being investigated. Next, we examined the extent of crypt labeling for p65276 and tried to correlate with changes in NF-κB activity and crypt hyperplasia utilizing tissue sections prepared from uninfected and days 5, 12 and 19 post-CR infected mouse distal colons from wild type C57Bl/6 and *Tlr4*2/2 mice. As depicted in Fig. 2J, p65276 labeling in sections from uninfected mouse distal was restricted to the base of the crypt. At days 5 and 12, p65276 labeling in the crypts of C57Bl/6 mice increased significantly (Fig. 2J) and correlated with changes in NF-κB activity recorded at these time points (see Fig. 2B) while p65276 labeling decreased significantly at day 19 (Fig. 2J). Interestingly, p65276 labeling at day 5 and particularly at days 12 and even 19 in *Tlr4*2/2 mice was significantly higher than the wild type counterpart and correlated with NF-κB activity at these time points (see Fig. 2F) and crypt hyperplasia (Fig. 1). Figure S1 further shows evidence of
Figure 2. TLR4-dependent and independent mechanism of NF-κB activation during TMCH. Relative levels of TLR4 in the cellular extracts prepared from the isolated crypts of uninfected normal (N) and CR infected (days 3–19) C57Bl/6 (A) and Tlr4^{-/-} (E) mice, respectively. Actin was used as internal control. Measurement of NF-κB activity in the colonic crypts of C57Bl/6 and TLR4^{-/-} mice. NFκB-p65 activity in the nuclear extracts prepared from the uninfected normal (N) and days 3–19 post infected C57Bl/6 (B) and Tlr4^{-/-} (F) mice was examined by utilizing TransAM NF-κB p65 Chemi Transcriptional Factor assay kit from Active Motif. *, p<0.05 vs. N; **, p<0.05 vs. day 5; †, p<0.05 vs. day 7; ‡, p<0.05 vs. day 12 (n = 3 independent experiments). Phosphorylation status of p65 subunit in the colonic crypts of C57Bl/6 and Tlr4^{-/-} mice. Relative levels of phosphorylated (p65^{Ser276}/p65^{Ser536}) and total p65 subunit in the cellular and nuclear extracts prepared form the uninfected normal (N) and days 3–19 post infected C57Bl/6 (C) and Tlr4^{-/-} (G) mice were determined by Western blotting with moiety-specific antibodies. Actin or LaminB were loading controls. CXCL-1/KC expression in the colonic crypts of C57Bl/6 and TLR4^{-/-} mice. Expression of CXCL-1/KC mRNA isolated from the crypts of uninfected normal (N) and days 3–19 post infected C57Bl/6 (D) and TLR4^{-/-} (H) mice respectively, was measured as readout for NF-κB activity, via real-time RT-PCR. *, p<0.05 vs. N; **, p<0.05 vs. day 12, (n = 3 independent experiments). I. Bacterial counts correlate with NF-κB activation in TLR4^{-/-} mice. Distal colonic homogenates, prepared from the uninfected and CR-infected (days 9–19) Tlr4^{-/-} mice were plated on McConkey agar and incubated at 37°C. Pink Citrobacter rodentium colonies were counted and plotted as log values (n = 3 independent experiments). J. Immunohistochemical staining of p65 subunit phosphorylated at Ser-276 in vivo. Paraffin embedded sections prepared from the distal colons of uninfected normal (N) and days 5, 12 and 19 post-infected wild type C57Bl/6 (WT) and Tlr4^{-/-} mice were stained with antibody specific for NFκB-p65 phosphorylated at Ser-276 and were analyzed with light microscopy. Percentages represent percent cells positive for p65^{Ser276} staining. (Scale bar = 75 μm; n = 3 independent experiments). doi:10.1371/journal.pone.0079432.g002
significant p65\textsuperscript{\texttt{276}} labeling at days 9, 12 and 19 compared to uninfected control which correlates with gross changes in the colonic mucosa. Thus, TLR4 requirement is not absolute for NF-\kappaB activation in response to CR infection.

Next, we examined the contribution of \beta-catenin towards crypt hyperplasia in response to CR infection in the wild type and \textit{ Tb4\textsuperscript{\texttt{*/-}}} mice, respectively. We began by measuring moiety-specific phosphorylation of \beta-catenin using specific antibodies. In the C57Bl/6 wild type mice, cellular and nuclear levels of Ser\textsuperscript{\texttt{45}}-phosphorylated \beta-catenin (\beta-Cat\textsuperscript{\texttt{45}}) increased between days 5–12 post-infection compared to uninfected control before declining at day 19 (Fig. 3Ai) and correlated with gross changes in hyperplasia. Increases in Ser\textsuperscript{\texttt{45}}-phosphorylation when normalized to total \beta-catenin levels however exhibited only subtle enhancement in phosphorylation intensity suggesting that increases in \beta-Cat\textsuperscript{\texttt{45}} may be due to a proportional increase in overall \beta-catenin abundance (Fig. 3Ai). Similarly, relative levels of \beta-catenin\textsuperscript{\texttt{552}} as well as de-phospho-\beta-catenin (dp\beta-Cat), both known to represent active \beta-catenin, increased significantly in both the cellular and nuclear extracts between days 5–12 compared to uninfected control before a rapid decline to basal levels at day 19 (Fig. 3Ai). As before, the increases in these species when compared to changes in total \beta-catenin were subtle. The alterations in various \beta-catenin species correlated with relative levels of downstream targets c-myc and cyclinD1 (Fig. 3Ai). In \textit{ Tb4\textsuperscript{\texttt{*/-}}} mice, increases in cellular and nuclear levels of \beta-Cat\textsuperscript{\texttt{45}} and \beta-Cat\textsuperscript{\texttt{552}} at days 5 and 12 compared to uninfected control (Fig. 3Bi), were mirror images of changes recorded in phosphorylated p65 at these time points (see Fig. 2G) while changes at other time points were less significant. In contrast, while the cellular levels of dp\beta-Cat followed the \beta-Cat\textsuperscript{\texttt{45}} and \beta-Cat\textsuperscript{\texttt{552}} pattern, nuclear levels (Fig. 3Bi) were similar to those recorded in the crypts of wild type C57Bl/6 mice (see Fig. 3Ai) except that the levels even at day 19 were significantly higher than uninfected control. The increases in these species were apparently a reflection of an overall increase in total \beta-catenin at these time points but unlike wild type crypts wherein, the total \beta-catenin returned to baseline at day 19, total \beta-catenin in \textit{ Tb4\textsuperscript{\texttt{*/-}}} crypts remained elevated than uninfected control (Fig. 3Bi). When we measured downstream targets, both c-myc and cyclinD1 exhibited kinetics similar to that recorded in wild type crypts except that the levels even at day 19 were still higher than uninfected control (Fig. 3Bi). These changes correlated well with persistence of crypt hyperplasia at this time point (see Fig. 1D). The changes in cellular/nuclear \beta-catenin and cyclinD1 in \textit{ Tb4\textsuperscript{\texttt{*/-}}} mice were further validated via immunohistochemistry. As depicted in Figure 3C, a significant increase in diffused cytosolic, nuclear and peri-nuclear staining for \beta-catenin was recorded at days 5 and 12 compared to uninfected control while nuclear staining declined at day 19. Similar staining pattern for cyclinD1 (Fig. 3D) coincided with Western blotting data (see Fig. 3Bi).

Cytosolic levels of free \beta-catenin are regulated by APC protein. We have shown previously that increases in cellular/nuclear levels of \beta-catenin in response to CR infection were neither due to mutational activation of \beta-catenin or loss of functional APC protein [35–37,43]. In the current study, relative increases in cellular levels of p312 were recorded at day 12 followed by a decline at day 19 in the colonic crypts of both the wild type and \textit{ Tb4\textsuperscript{\texttt{*/-}}} mice suggesting that increases in \beta-catenin at day 12 were not due to loss of wild type APC (Fig. 3E). At the same time and as shown by us previously [35], increases in \beta-catenin were probably due to GSK-3\beta's phosphorylation at Ser-9 which results in its inactivation. These changes in phospho-GSK-3\beta were more pronounced in the colonic crypts of \textit{ Tb4\textsuperscript{\texttt{*/-}}} mice compared to the wild type counterpart (Fig. 3E). Finally, consistent with the earlier finding [44], we observed significant increases in PKC\zeta, the kinase that phosphorylates Ser-9 of GSK3\beta [35], at day 12 post-infection compared to uninfected control in the colonic crypts of \textit{ Tb4\textsuperscript{\texttt{*/-}}} mice suggesting (but certainly not proving) PKC\zeta-mediated inactivation of GSK-3\beta at this time point while changes in wild type mice were less significant (Fig. 3F).

Differential Regulation of \beta-catenin and NF-\kappaB Pathways in vitro

So far, we have seen that both increases in NF-\kappaB and \beta-catenin signaling following CR infection coincide with colonic crypt hyperplasia in wild type as well as \textit{ Tb4\textsuperscript{\texttt{*/-}}} mice, respectively. To investigate if an interplay between the two pathways exists, we opted for \textit{ in vitro} studies to focus on two proteins, Wnt2b and Wnt5a that activate Wnt/\beta-catenin and NF-\kappaB pathways, respectively [43,27]. Semi-quantitative PCR revealed significant increases in both Wnt2b and Wnt5a expression in YAMC cells at 48 h post-CR infection compared to uninfected control (Fig. 4A). During measurement of \beta-catenin/Tcl4-dependent TOP-flah reporter activity, CR infection induced a 2-fold increase in reporter activity at 48 h post-infection (Fig. 4B). In response to siRNA-mediated Wnt2b knockdown, reporter activity decreased significantly and was not rescued with CR infection (Fig. 4B). Wnt5a knockdown on the other hand, resulted in an increase in reporter activity which was further enhanced following CR infection (Fig. 4B). Next we examined the effect of adding purified Wnt2b and Wnt5a proteins on reporter activities. CR infection compared to uninfected control increased the reporter activity by ~2-fold as described before (Fig. 4C). Si-RNA to Wnt2b almost completely attenuated the reporter activity while addition of purified Wnt5a did not rescue the inhibitory effects (Fig. 4C). Si-RNA to Wnt5a had no effect on the CR-induced reporter activity which remained elevated while addition of Wnt2b further enhanced the reporter activity (Fig. 4C). To confirm if these changes were mediated by increases in \beta-catenin, we stained YAMC cells for \beta-catenin. Immunofluorescent studies revealed predominantly membranous staining for \beta-catenin in uninfected cells (Fig. 4D). In contrast, significant nuclear labeling of \beta-catenin was observed in response to CR infection (Fig. 4D). Interestingly, siRNA to Wnt2b almost completely blocked \beta-catenin nuclear translocation while siRNA to Wnt5a had no effect (Fig. 4D). Similarly, during scratch-induced wound assay in the YAMC cells, knockdown of Wnt2b but not Wnt5a significantly blocked \beta-catenin-dependent wound healing (Fig. 4E). Figure 4F is a representative bar graph showing percent migration at 16 h. Thus, Wnt2b and not Wnt5a promotes Wnt signaling in response to CR infection. We next examined the effect of siRNA-mediated knockdown of Wnt2b and Wnt5a on NF-\kappaB activity \textit{ in vitro}. During the DNA binding assay, NF-\kappaB activity increased significantly in response to CR infection compared to uninfected control while siRNA to Wnt2b had no effect (Fig. 4G). On the other hand, siRNA to Wnt5a almost completely attenuated CR-induced NF-\kappaB activity (Fig. 4G). Thus, Wnt5a and not Wnt2b regulates NF-\kappaB activity in response to CR infection.

As proof-of-principle, we further characterized the role(s) of purified Wnt2b and Wnt5a proteins in modulating the Wnt and NF-\kappaB pathways \textit{ in vitro}. During immunostaining, \beta-catenin remained membrane bound in uninfected cells while a predominant nuclear staining was recorded in CR infected cells (Fig. 5A), as described before (see Fig. 4D). Interestingly, purified Wnt2b was more potent than even CR infection in facilitating \beta-catenin’s nuclear import while a combination of Wnt2b+CR did not necessarily yield an additive response (Fig. 5A). Addition of purified Wnt5a on the other hand, did not affect either \beta-catenin
Figure 3. Effect of CR infection on Wnt/\(\beta\)-catenin signaling in vivo. Relative levels of phosphorylated (\(\beta\)-Cat\(^{45}\)/\(\beta\)-Cat\(^{552}\)), de-phosphorylated (\(d\beta\)-Catenin) and total \(\beta\)-catenin in the cellular and nuclear extracts prepared from the uninfected normal (N) and days 3–19 post infected wild type (WT) C57Bl/6 (Ai) and Tlr4\(^{-/-}\) (Bi) mice were determined by Western blotting with moiety-specific antibodies. Actin or LaminB were loading controls. Downstream targets of Wnt/\(\beta\)-catenin signaling are upregulated in response to CR infection. Relative levels of cyclinD1 and c-Myc in the nuclear extracts prepared from the uninfected normal (N) and days 3–19 post infected wild type (WT) C57Bl/6 (Aii) and Tlr4\(^{-/-}\) (Bii) mice were determined by Western blotting. LaminB was used as loading control. C and D. Immuno-staining for \(\beta\)-catenin and cyclinD1 in Tlr4\(^{-/-}\) mice. Paraffin embedded sections prepared from the distal colons of uninfected normal (N) or 5, 12 and 19-days post-CR infected Tlr4\(^{-/-}\) mice were stained for \(\beta\)-catenin (C) or cyclinD1 (D).
levels or nuclear translocation while a combination of Wnt5a+CR restored β-catenin’s nuclear localization (Fig. 5A). During a time course with purified proteins, addition of Wnt2b led to significant increases in relative levels of nuclear β-catenin at 24 hr and the levels persisted until 72 hr post-addition (Fig. 5Bi). These increases were either at par or even higher than those recorded with CR infection (Fig. 5Bi). Interestingly, we did not observe any detectable levels of either p65 phosphorylated at Ser-536 (p65S536) or β-catenin’s downstream targets Slug and Snail following addition of purified Wnt2b while CR infection alone promoted increases in these proteins at indicated times (Fig. 5Bi). Wnt5a addition on the other hand, led to significant increases in relative levels of nuclear p65S536 at 24 hr which persisted until 48 hr before declining at 72 hr similar to those recorded with CR infection alone while Wnt5a only had nominal effect on β-catenin levels at these time points (Fig. 5Bi). Similar to Wnt2b, Wnt5a did not affect Slug and Snail expression at either time point (Fig. 5Bi). Next, during measurement of NF-κB activity, CR infection compared to uninfected control, caused a reproducible increase in NF-κB activity at 24 hr which persisted until 72 hr (Fig. 5C). Following addition of Wnt5a, a significant increase in NF-κB activity was recorded at 24 hr which persisted until 72 hr while addition of Wnt2b had subtle effect on NF-κB activity (Fig. 5C). Finally, based on our previous report regarding activation of MEK/ERK pathway following CR infection in the colonic crypts [41], we performed scratch-induced wound assay in YAMC cells in the presence or absence of MEK inhibitor PD98059 (MEKi) to understand if MAPK signaling may be contributing towards CR+Wnt2b-induced wound healing. As depicted in Figure 5D, CR infection compared to uninfected cells, caused almost complete wound closure between 12–16 h (Fig. 5D). Treatment of uninfected cells with MEKi only caused a subtle failure in cell migration while CR infection of MEKi-treated cells neutralized the inhibitory effect (Fig. 5D). Si-RNA to Wnt2b significantly blocked CR-induced cell migration (Fig. 5D) as described earlier (see Fig. 4E). Interestingly, a combination of siRNA-Wnt2b+MEKi almost completely abrogated CR-induced cell migration (Fig. 5D) suggesting that Wnt2b/MEK cross-talk may be required for CR-induced cell migration. Figure S2A is a representative bar graph showing percent migration at 16 h. Taken together, these studies clearly implicate both canonical and non-canonical Wnt pathways in the differential regulation of β-catenin and NF-κB in response to CR infection.

Blocking β-catenin in vivo Abrogates Crypt Hyperplasia and Reduces Tumorigenesis

Our next aim was to understand the interplay between the β-catenin and NF-κB pathways in vivo in response to CR infection in the wild type (WT) mice and mice deficient for TLR4. In response to either CR infection or CR infection+vehicle treatment of either strain, significant increases in both active and total β-catenin was observed in the crypt cellular and nuclear extracts along with increases in its downstream target cyclinD1, compared to uninfected control (Fig. 6A, B). When WT or Tlr4−/− mice were treated with nanoparticle-encapsulated siRNA to β-catenin (si-β-Cat), we observed almost complete loss of nuclear β-catenin with concomitant decreases in cyclinD1 (Fig. 6A, B). As expected, si-β-Cat treatment had no effect on active (p65S276, p65S327) or total NF-κB-p65 which remained elevated in CR infected crypt cellular and nuclear extracts (Fig. 6A, B). Interestingly, si-β-Cat treatment significantly blocked β-Cat522 nuclear staining concomitant with decreases in cell proliferation as it was determined by reduced Ki-67 staining in treated samples without grossly affecting the mucosa (Fig. 6C, D). To see if downstream targets of Wnt/β-catenin signaling and putative markers of stem cells were affected, we next examined the effect of si-β-Cat treatment on CD44 and Dclk1 staining in the two sets of mice. In response to either CR infection or CR infection+vehicle treatment, staining for both CD44 and Dclk1 increased in the colonic crypts (Fig. 6E, F). Following si-β-Cat treatment, staining for both CD44 and Dclk1 were reduced to baseline (Fig. 6E, F). Interestingly, the effect of si-β-Cat treatment on CD44 was much more pronounced in sections prepared from Tlr4−/− mice compared to wild type counterpart (Fig. 6F). We have shown recently that functional cross-talk between Notch and NF-κB pathways regulate crypt hyperplasia and/or tumorigenesis in response to CR infection [38]. We therefore examined the effect of si-β-Cat treatment on cancer stem cell markers in ApcMins/+ mice to see if tumorigenesis is affected. si-β-Cat treatment to ApcMins/+ mice significantly attenuated increases in β-Cat522, CD44, Dclk1 and CD133 that halted the growth of mutated crypts (Fig. 7) without affecting NF-κB-p65S536 signaling (data not shown). Thus, β-catenin and not necessarily NF-κB regulates crypt hyperplasia and tumorigenesis in response to CR infection.

Evidence of β-catenin and not NF-κB-induced Crypt Hyperplasia in CAST.11M Mice

Castaneus mice contain a CAST/Ei region on chromosome 11 (CAST.11M) on an otherwise C57BL/6 genetic background. Genetic variation within an 83 Mb region on mouse chromosome 11 controls host response to anthrax lethal toxin and resistance to infection by the Sterne strain of Bacillus anthracis [46,47]. We compared these mice with wild type C57BL/6 to determine if differences exist in the extent of β-catenin and NF-κB activation and associated crypt hyperplasia in these mice in response to CR infection. Following CR infection, C57BL/6 mice exhibited significant crypt hyperplasia as revealed by Ki-67 staining at day 9 which peaked by day 12 and plateaued by day 19 (Fig. 8Ai). In CAST.11M mice however, Ki-67 staining and crypt lengths at day 12 and particularly at day 19 compared to uninfected controls were 1.5–2 fold higher than those recorded in C57BL/6 mice at the same time points (Fig. 8Aii). Transmission electron microscopy revealed no difference in bacterial binding at the luminal surface in either strain (Fig. 8B); yet, C57BL/6 mice had significant disruption of the tight junctions (Fig. 8B), an elevated serum FITC-Dextran levels (Fig. 8B) and recruitment of CD3+ T-cells and F4/80+ macrophages to the sub-epithelial regions compared to CAST.11M mice (Fig. 8S). Mechanistically, in C57BL/6 mice, Ki-67 kinetics coincided with increases in nuclear accumulation of β-catenin and its downstream target cyclinD1 at days 9 and 12 compared to uninfected control while the levels declined at day 19 (Fig. 8Ci, Cii). In addition, p65S276/p65S327 subunit expression (Fig. 8Dii) and NF-κB activity (Fig. 8Dii) increased significantly at days 9 and 12 with declining trend at day 19 that correlated with downstream target CXCL-1 expression (Fig. 8Ei). In CAST.11M mice on the other hand, we observed a delayed response wherein, nuclear β-catenin levels did not go up until day...
Figure 4. Evidence of β-catenin and NF-κB interplay in vitro. A. Measurement of Wnt2b and Wnt5a expression in YAMC (Young Adult Mouse Colon) cells in vitro. YAMC cells (5 × 10^5) were either uninfected (N) or infected with CR at 90:1 MOI for 3 hr. Cells were washed thoroughly to remove bacteria and incubated in fresh medium containing antibiotics for indicated period of time. Total RNA was examined for the expression of Wnt2b and Wnt5a via RT-PCR. GAPDH was used as loading control. B. Effect of Wnt2b knockdown on reporter activity. YAMC cells were transiently transfected with TOPflash plasmid and with siRNA specific to Wnt2b and Wnt5a, respectively. After 24 h, cells were infected with CR at 90:1 MOI for 3 h, washed to remove bacteria followed by measurement of reporter activity at 48 h using Renilla luciferase as internal control (*, p < 0.05 vs. N; **, p < 0.05 vs. CR; *, p < 0.05 vs. CR; n = 3 independent experiments). C. Effect of Wnt2b and Wnt5a addition on reporter activity. YAMC cells, following transient transfection for 24 h with TOPflash plasmid and with siRNA specific to Wnt2b and Wnt5a, respectively. After 24 h, cells were treated with purified Wnt2b or Wnt5a, infected with CR at 90:1 MOI for 3 h, washed to remove bacteria followed by measurement of reporter activity at 48 h using Renilla luciferase as internal control (*, p < 0.05 vs. CR; **, p < 0.05 vs. CR; n = 3 independent experiments). D. β-catenin expression in YAMC cells in vitro. YAMC cells (5 × 10^5) were either uninfected (N) or infected with CR at 90:1 MOI for 3 hr. Cells were washed thoroughly to remove bacteria and incubated in fresh medium containing antibiotics for indicated period of time. Total RNA was examined for the expression of β-catenin via RT-PCR. GAPDH was used as loading control. E. Effect of Wnt2b and Wnt5a addition on β-catenin expression. YAMC cells, following transient transfection for 24 h with TOPflash plasmid and with siRNA specific to Wnt2b and Wnt5a, respectively. After 24 h, cells were treated with purified Wnt2b or Wnt5a, infected with CR at 90:1 MOI for 3 h, washed to remove bacteria followed by measurement of β-catenin expression at 48 h using western blot analysis. F. Effect of Wnt2b and Wnt5a addition on cell migration. YAMC cells, following transient transfection for 24 h with TOPflash plasmid and with siRNA specific to Wnt2b and Wnt5a, respectively. After 24 h, cells were treated with purified Wnt2b or Wnt5a, infected with CR at 90:1 MOI for 3 h, washed to remove bacteria followed by measurement of cell migration at 48 h using Transwell assay. G. Effect of Wnt2b and Wnt5a addition on NF-κB activity. YAMC cells, following transient transfection for 24 h with TOPflash plasmid and with siRNA specific to Wnt2b and Wnt5a, respectively. After 24 h, cells were treated with purified Wnt2b or Wnt5a, infected with CR at 90:1 MOI for 3 h, washed to remove bacteria followed by measurement of NF-κB activity at 48 h using ELISA assay.
12 with sustained expression at day 19 (Fig. 8Gi) which coincided with cyclinD1 expression (Fig. 8Gii) and Ki-67 kinetics at these time points. Interestingly, CAST.11M mice did not exhibit any significant increase in phosphorylated/total p65 subunit expression (Fig. 8Di), NF-κB activity (Fig. 8Dii), CXCL-1 expression (Fig. 8Ei) or serum FITC-Dextran levels (Suppl. Fig. 2) at these time points despite adequate bacterial binding to the colonic mucosa (Fig. 8B). Thus, β-catenin can regulate crypt hyperplasia in response to CR infection in the absence of NF-κB signaling.

Discussion

Wnt/β-catenin and NF-κB are independent pathways involved in the regulation of physiological and pathological effects related to the development, immune function, inflammation, tumorigenesis and frank malignancy. The recent discovery of functional cross-regulation between the two pathways has established complex roles for Wnt/β-catenin and NF-κB signaling in the pathogenesis of various diseases including cancer. Indeed, the mechanisms and biological effects of the cross-regulation of the two pathways remains an area of intense investigation. It is now evident that strategies targeting the cross-regulation between these pathways may be a promising direction for future cancer therapeutics.

Recognition of Gram negative bacterial products such as lipopolysaccharide (LPS) by toll like receptor-4 (TLR4) and subsequent activation of NF-κB not only activates host defense against pathogenic bacteria, but can also provide mucosal protection. In pathological situations however, TLR4 is upregulated in both Crohn’s disease and ulcerative colitis, two components of human IBD, leading to pro-inflammatory signaling via NF-κB [42,48]. We recently showed that NF-κB activation in response to CR infection was dependent upon signaling via TLR4 [41]. In this study, we observed significant increases in cellular levels of TLR4 in the colonic crypts of CR infected mice suggesting critical role(s) for TLR4 in transducing signals in response to CR infection. To definitively implicate TLR4 in CR-induced NF-κB activation, we utilized Tbr4-/- mice in the B6 background wherein, initial kinetics of NF-κB activation at day 3 highlighted the need for the presence of TLR4; NF-κB activation at day 5 however, may represent a homeostatic response to counter bacterial infection while NF-κB activation at day 12 which correlated with peak hyperplasia (see Fig. 1) may represent an intrinsic pathway activation independent of TLR4. This is in contrast to decreased epithelial proliferation and significant epithelial injury recorded during DSS-induced colitis in Tbr4-/- mice [42]. These differences may be attributable to the nature of the colitis-inducing agent: CR infection may represent more dramatic changes in β-catenin cellular abundance and downstream target expression in Tbr4-/- mice and these changes directly correlated with increases in crypt hyperplasia in infected mice. A definitive role for β-catenin in CR-induced crypt hyperplasia in Tbr4-/- mice was established by blocking increases in β-catenin in vivo which resulted in attenuation of the hyperplastic response. These results clearly contradict the notion that TLR4 is somehow required for activation of the Wnt/β-catenin pathway. Indeed, a recent study by Sodhi CP et al. [28] suggested that TLR4 actually inhibits β-catenin and impairs enterocyte proliferation in the small intestine of neonatal mice with experimental necrotizing enterocolitis (NEC). This effect however, was not seen in the colon of the same neonatal mice nor was it found anywhere in the intestine of adult mice. Thus, even though our findings particularly in Tbr4-deficient mice corroborate with those described during NEC, the regulation of Wnt/β-catenin signaling via TLR4 seems more complex than what was initially envisioned.

Continuing with our theme of interplay between the Wnt/β-catenin and the NF-κB pathways, an elegant study recently reported that constitutive IKKβ/NF-κB activation synergized with Wnt signaling to promote intestinal tumorigenesis.
Figure 5. Evidence of differential β-catenin and NF-κB signaling by Wnt ligands in vitro. A. YAMC cells were either treated with control media or infected with CR (90:1 MOI) for 3 h followed by washing to remove bacteria. Uninfected or CR infected cells were incubated with purified β-Catenin/NF-κB Interplay and Disease Pathogenesis

B. Representative images of β-catenin and NF-κB localization in YAMC cells infected with CR and treated with Wnt ligands. DAPI staining is used as a nuclear marker.

C. Graphical representation of NF-κB activity (OD at 560 nm) over time for different conditions.

D. Time-lapse imaging showing the progression of β-catenin and NF-κB localization in YAMC cells infected with CR and treated with Wnt ligands at 0h, 12h, and 16h.
Wnt2b or Wnt5a for 24 hr followed by immuno-staining for β-catenin. Nuclei were stained with DAPI (n = 3 independent experiments). B. Effect of Wnt2b and Wnt5a addition on cellular levels of β-catenin and NF-κB. YAMC cells described in A were incubated with purified Wnt2b or Wnt5a for 24, 48 and 72 hr followed by Western blotting with antibodies for; p65 subunit phosphorylated at Ser-536 (p65 536), β-catenin, Slug and Snail, respectively. Actin was used as loading control. C. Effect of Wnt2b and Wnt5a addition on NF-κB activity. YAMC cells described in B were subjected to nuclear extraction followed by measurement of NF-κB activity utilizing TransAM NF-κB p65 Chemi Transcriptional Factor assay kit from Active Motif (*, p < 0.05 vs. N; n = 3 independent experiments). D. Effect of MEK inhibitor (MEKi) on wound healing. YAMC cells were transfected with control or Wnt2b-specific siRNA for 24 h followed by incubation with MEKi. At 48 h, cells were wounded with a plastic pipette tip and cell migration was followed for 12–16 h (n = 3 independent experiments).

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[52]. We showed previously that functional cross-talk between β-catenin and NF-κB is integral to the hyperproliferative effects of progastrin on proximal colonic crypts [53]. In the current study however, we have discovered that despite significant activation of the NF-κB pathway in the distal colons of wild type or Tlr4−/− mice or in ApcMin/+ mice, knockdown of β-catenin significantly impacted both hyperplasia and tumorigenesis in response to CR infection. Since a combination of canonical and non-canonical Wnt signaling regulates the interplay between the Wnt/β-catenin and NF-κB pathways [54], we have further elucidated the roles of two Wnt ligands, Wnt2b and Wnt5a in the differential regulation of these pathways in vitro. Based on functional assays with purified proteins, we conclude that Wnt2b but not Wnt5a regulates the Wnt/β-catenin pathway while Wnt5a and not Wnt2b is involved in the activation of the NF-κB pathway, respectively. In previous studies, Wnt2b is shown to be transiently expressed in the primitive streak during gastrulation and has been suggested to primarily play a supportive role in gastrulation and organogenesis [55]. It is upregulated in liver and biliary epithelial cells of patients with primary biliary cirrhosis [56], and in colonic mucosa of patients with inflammatory bowel disease [57]. Wnt2b is also shown to be expressed in several types of human cancer, such as basal cell carcinoma, gastric cancer, breast cancer, head/neck squamous cell carcinoma, cervical cancer and leukemia. More recently, canonical Wnt2/2b signaling is shown to be required for specification of the lung endoderm progenitors in the developing foregut [58]. Similarly, Wnt5a activates both canonical and non-canonical Wnt signaling pathways and is implicated in a variety of cellular processes during development and carcinogenesis. Importantly, the expression of Wnt5a protein is controlled by the NF-κB signaling pathway, which may be implicated as an essential mediator not only for infection, but also for cancer development [54]. While we present evidence of differential β-catenin and NF-κB regulation by the two Wnt ligands in primary colonicocytes in vitro, studies in new mouse models allowing specific depletion of genes encoding these proteins in the epithelial cells of the colon will be required to address the potential role(s) of these ligands in either crypt hyperplasia or tumorigenesis in response to CR infection.

The dominance of β-catenin-induced crypt proliferation in response to CR infection was further validated in a Castaneus strain (CAST;11M) that exhibits high susceptibility to Anthrax’s lethal toxin (LT) [47] due to a region at chromosome 11 that encodes the LT-responsive CAST/Ei allele of Nrp1b inflammasome [47]. Interestingly, LT-mediated lethality in mice was found to be independent of TLR4 function [46]. Thus, while it is intriguing to observe a muted NF-κB response in the distal colons of these mice, it is not known whether modifier genes at chromosome 11 are necessarily interfering with NF-κB activation in response to CR infection. Whatever the case may be, the fact that we still observed significant increases in β-catenin in these mice that correlated with crypt hyperplasia further proves our hypothesis that β-catenin and not necessarily NF-κB regulates crypt hyperplasia in response to CR infection. Given that deregulation of the components of the Wnt/β-catenin signaling pathway has been implicated in a wide spectrum of diseases including cancer, our studies further provide a rationale for targeting the Wnt/β-catenin pathway to treat diseases with infectious etiology.

Materials and Methods

Mice, CR Infection and Induction of Hyperplasia

This study was carried out in strict accordance with the recommendations in the Guide for the Care and Use of Laboratory Animals of the National Institutes of Health. All protocols were approved by University of Kansas Medical Center Animal Care and Use Committee. All efforts were made to minimize suffering. Male Helicobacter pylori-free ApcMin/+ and Tlr4−/− mice were procured from Jackson Laboratory, Bar Harbor, Maine, USA. B6.CAST.11M Castaneus mice were kindly provided by Drs. Aldons J. Luis and Richard C. Davis (University of California at Los Angeles) and the generation of these mice has been described previously [59]. B6.CAST.11M mice contain a CAST/Ei region in chromosome 11 (~31.5 Mb to the terminus) on an otherwise C57BL/6J genetic background [59,60]. All the mice were maintained in a specific pathogen-free (including helicobacter and parvovirus) environment, and generally used between 5 and 6 weeks of age. As control groups, either littermates or WT C57Bl/6J mice of identical background were used. Transmissible murine colonic hyperplasia was induced in ApcMin/+; Tlr4−/− and wild type C57Bl/6J mice by oral inoculation with a 16-h culture of CR (biotype 4280, ATCC, 10⁶CFUs) identified as pink colonies on MacConkey agar, as previously described [34–39,41,43,44,61–67]. Age- and sex-matched control mice received sterile culture medium only.

RNA Extraction, Semi-quantitative and Real-time PCR

Total RNA was extracted from isolated colonic crypts by using TRI Reagent. Following cDNA synthesis using Superscript II and random primers, gene products specific to Wnt-2b (F: 5'-CACCCGACGTGATCTTGTCT-3'; R: 5'-TGGTTCCTGTCA-CTCCCTGTCAAC-3') and Wnt5a (5'-CTGCGGAGCTTTCTC-CAAGG-3'; 5'-CTCTAGGCTCCAGAACTCC-3') were identified by performing semi-quantitative PCR using 1/20 dilution of the cDNA. For each gene product, the amplification cycle number was chosen empirically within the linear range. The PCR products were separated by polyacrylamide gel electrophoresis and visualized by ethidium bromide staining of the gels under UV light. Gel data were recorded with the Bio-Rad FluorS Imaging System, and relative densities of the bands were determined with Quantity One software (Bio-Rad, Hercules, CA). Gene expression was normalized with β-actin expression. Total RNA samples were also subjected to real-time polymerase chain reaction (PCR) by SYBR chemistry (SYBR Green I; Molecular Probes, Eugene, OR) using primers specific for CXCL-1/KC (F: 5'-GCCAATGAGCT-TGGGCTGTCAATGC-3'; R: 5'-CTGCGGAGCTTTCTGAGC-3') and GAPDH (F: 5'-AATTTGGGATGTGGAAAG-3'; R: 5'-ACACATTGGGTTAGGAACA-3'). The changes in mRNA were expressed as fold change relative to control with ± SEM value.
Figure 6. Signaling via β-catenin is integral to crypt hyperplasia. CR-infected wild type C57Bl/6 (A) and Tlr4−/− (B) mice either received vehicle (V) or ip injections of nanoparticle-encapsulated β-catenin siRNA (Si) every alternate day for 10 days. Colonic crypts were isolated and subjected to cellular or nuclear extract fractionation followed by Western blotting for indicated proteins. Actin and LaminB were used as loading controls. C–F. Paraffin embedded sections prepared from the distal colons of uninfected normal (N), CR infected (CR), CR infected+vehicle treated (V) or CR infected+β-catenin siRNA treated (si-Cat) mice were stained with: C and D. H&E for gross morphology (upper panel), β-catenin phosphorylated at Ser-552 to detect active β-catenin (middle panel) and Ki-67 for proliferation (lower panel); E and F. CD44 (upper panel) and Dclk1 (lower panel) were also stained to determine the effect of β-catenin knockdown on stem cells. Percentages represent percent cells positive for indicated markers. Scale bar: 125 μm; n = 2 independent experiments.

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DNA Binding Assay

For DNA binding assay, nuclear extracts were prepared from the distal colonic crypts of uninfected, or days-3, 5, 7, 9, 12 and 19 infected mice, respectively. The young adult mouse colon (YAMC) cells were maintained in RPMI 1640 medium supplemented with 10% fetal bovine serum, 2 mM glutamine, 50 μg/ml gentamicin, 100 units/ml penicillin, 100 μg/ml streptomycin, and 5 units/ml IFNγ in a humidified incubator with 5% CO2 at 33°C. YAMC cells (5 x 10⁵) were transfected with either 100 nmol/L of scrambled siRNA or siRNAs specific for Wnt2b (sc-155356) and Wnt5a (sc-41113) using 2–8 μl of transfection reagent (sc-29528; Santa Cruz Biotechnology Inc., Santa Cruz, CA) for 24 h. CR strain DBS 100 (ATCC cat. # 51459™) was grown under aerophilic conditions on Luria-Bertani (LB) agar plates for 24 h at 37°C and cultured in LB broth O/N at 37°C. RPMI, containing 0.45% glucose was inoculated with a 1:20 dilution of a standard LB overnight culture and incubated for 2 h at 37°C. Monolayers of 5 x 10⁵ YAMC cells at ~50% confluence or cells transfected with various siRNAs after 24 h were infected with CR at a multiplicity of infection (MOI) of 90 or the medium alone (as a control) or incubated with purified Wnt2b and Wnt5a (@250 ng/ml) for 3 h at 37°C in 5% CO₂. After 3 h, medium was changed and replaced with fresh medium plus antibiotics to ensure complete absence of live bacteria. Relative levels of activated p65-NF-κB in the nuclear extracts were measured using the

![Figure 7. Effect of β-catenin knockdown on tumorigenesis in Apc<sup>Mim1</sup> mice.](image)

Representative photomicrographs of paraffin embedded sections prepared from the distal colons of uninfected normal (N), CR-infected at day 12, CR-infected mice at 3 months of age with tumor (CR<sub>T</sub>) and CR<sub>T</sub> mice treated with nanoparticle-encapsulated β-catenin siRNA (si-Cat<sub>T</sub>). Sections were stained with antibodies for: β-catenin phosphorylated at Ser-552 (β-catenin<sup>552</sup>), and cancer stem cell markers CD44, Dclk1 and CD133, respectively. Percentages represent percent cells positive for indicated markers. Scale bar: 50 μm; n = 2 independent experiments.

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Figure 8. B6.CAST.11M mice exhibit a hyperplastic but a muted inflammatory response to CR infection. A. Representative photomicrographs of paraffin embedded sections prepared from the distal colons of uninfected normal (N) and days 9–19 post-CR infected C57Bl/6 (upper panel) or CAST.11M (lower panel) mice and stained for Ki-67 (percentages represent percent cells positive for Ki-67). Scale bar: 50 μm; n = 3 independent experiments. B. Electron Microscopy. Distal colonic fragments from CR infected wild type C57Bl/6 (WT) or CAST.11M mice were subjected to transmission electron microscopy to detect disruption in tight junctions (arrows). Two representative images are shown for each group of mice. C. Relative levels of phosphorylated (β-Cat552), de-phosphorylated (dpβCatenin) and total β-catenin (Ci) and cyclinD1 (Cii) in the nuclear
extracts prepared form the uninfected normal (N) and days 9, 12 and 19 post infected wild type C57Bl/6 and B6.CAST.11M mice, were determined by Western blotting with moiety-specific antibodies. LaminB was used as loading control. DI. Nuclear extracts prepared from the distal colonic crypts of groups of p6559, p6556 and total p65, respectively. LaminB was used as loading control. DIi. Measurement of NF-κB activity. NF-κB-p65 activity in the nuclear extracts prepared from the groups of mice described in C was measured by utilizing TransAM p65 Chemi Transcriptional Factor assay kit from Active Motif. E. CXCL-1/KC expression in the colonic crypts. Expression of CXCL-1/KC mRNA isolated from the distal colonic crypts of C57Bl/6 (Ei) or CAST.11M (Eii) mice was measured as readout for NF-κB activity, via real-time RT-PCR. *

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TransAM p65 NF-κB Chemi Transcription Factor Assay Kit from Active Motif (Carlsbad, CA) as per manufacturer’s instructions.

Luciferase Reporter Gene Assay

Transfection experiments were carried out using Lipofectamine 2000 reagent (Invitrogen Life Technologies), according to the manufacturer’s instructions. YAMC cells were maintained as described above. For transient transfections, cells were: i) co-transfected with Tcf-4 reporter plasmid (TOPFlash; TOP) or a mutant Tcf-binding site (FOPFlash; FOP) along with pRL-TK Renilla vector as internal control, or ii) transfected with either 100 nmol/L of scrambled siRNA or siRNAs specific for Wnt2b (sc-155536) and Wnt5a (sc-41113) using 2–8 μl of transfection reagent (sc-29328; Santa Cruz Biotechnology Inc., Santa Cruz, CA). Twenty four hour post-transfection, cells were either uninfected (medium only) or infected with CR (@1:90 MOI) or incubated with purified Wnt2b and Wnt5a (@250 ng/ml) at 37°C in 5% CO₂ for 3 hr followed by washing to remove bacteria. At indicated times, cells were processed for measurement of luciferase activity with a luminometer and the values were normalized to the internal control. All transfection experiments were repeated at least three times.

Knockdown of Wnt/β-catenin Signaling in vivo and Western Blotting

For blocking β-catenin in vivo, we injected intraperitoneally, β-catenin siRNA incorporated into poly (lactide-co-glycolide) acid nanoparticles (siβ-Cat-NPs) (at 4 μM/Kg. body wt.) every alternate day for 10-days followed by euthanasia at 12-days post-infection [61,68]. These PLGA-NPs were synthesized using a double emulsion solvent evaporation technique as described [68]. Distal colonic crypts were isolated at various time points as described [34–39,53,61,67] and total crypt cellular or nuclear extracts (30–50 μg protein/lane), were subjected to SDS-PAGE and electrotransferred to nitrocellulose membrane. The membranes were blocked with 5% BSA or 5% nonfat dried milk. Membranes were incubated either 100 μmol/L of scrambled siRNA or siRNAs specific for Wnt2b (sc-155536) and Wnt5a (sc-41113) using 2–8 μl of transfection reagent (sc-29328; Santa Cruz Biotechnology Inc., Santa Cruz, CA). Twenty four hour post-transfection, cells were either uninfected (medium only) or infected with CR (@1:90 MOI) or incubated with purified Wnt2b and Wnt5a (@250 ng/ml) at 37°C in 5% CO₂ for 3 hr followed by washing to remove bacteria. At indicated times, cells were processed for measurement of luciferase activity with a luminometer and the values were normalized to the internal control. All transfection experiments were repeated at least three times.

Histology, Immunofluorescence (IMF) and Immunohistochemistry (IHC)

For histology, tissues were fixed with 10% neutral buffered formalin or in Carnoy’s fixative (60% methanol, 30% chloroform, and 10% acetic acid) prior to paraffin embedding. Paraffin-embedded sections (5 μm) were stained with H&E for gross morphology and the pictures were obtained with an Eclipse E1000 microscope (Nikon). IMF to detect β-catenin in YAMC cells treated as described above or IHCs to detect Ki-67, p65527, β-catenin, cyclinD1, CD44, CD133 and Dclk1 were performed on 5-μm-thick paraffin embedded sections prepared from the distal colons of uninfected normal, 3, 5, 7, 9, 12 and 19 days CR-infected or CR-infected and siβ-Cat-NPs treated mice utilizing Envision+System-HRP (DAB; DakoCytomation, Carpinteria, CA) with microwave accentuation as described previously [34,36–39,53,61,67]. Antibody controls included either omission of the primary antibody or detection of endogenous IgG staining with goat anti-mouse or anti-rabbit IgG (Calbiochem, San Diego, CA). The visualization was carried out using light microscopy. IHC data were graded based on percent of stained cells showing a range of immunoreactivity of: <10% of crypt epithelial cells stained positive, 10 to 30% positivity, 30 to 60% positivity or >60% positivity. A minimum of either ~200 crypts or ~2000 cells/field were counted for each experimental group. For statistical analysis of the staining scores, Student’s t-test was performed as described below.

Wound Healing Assay and Electron Microscopy

YAMC cells were transfected with siRNA specific to Wnt2b and Wnt5a, respectively for 24 h. Cells were subsequently treated with control media or with CR (@90:1 MOI) for 3 h followed by washing to remove bacteria. At 48 h, cells were wounded with a plastic pipette tip. After removing debris and adding fresh media, cell migration was followed for 12-16 h [61]. For electron microscopy, samples of the distal colon from uninfected normal or 9, 12 and 19 days’ CR-infected mice were minced into small cubes and fixed in 4% paraformaldehyde and 2% glutaraldehyde in Cacodylate buffer (0.1 M Sodium Cacodylate, pH 7.6) overnight at room temperature and postfixed in 1% Osmium tetroxide for 90 minutes. The fixed tissues were dehydrated through a graded series of ethanols and embedded in epon-araldite resin and maintained for 48 hours at 60°C to polymerize. Ultra-thin (100 nm) sections cut on a Leica UC-6 ultramicrotome were put on glow discharged 300 mesh copper grids and stained with Uranyl Acetate and Sato’s Lead to enhance contrast. Ultra-thin sections were examined with a Hitachi H-7600 electron microscope.

Statistical Analysis

Experiments were repeated three times with consistent results. Data were expressed as mean values ± standard error. Statistical analyses for all studies were performed using unpaired, two-tailed Student’s t-tests and one-way analysis of variance (ANOVA) for multiple group comparisons (GraphPad Prism 5, San Diego, CA). *p-values <0.05 were considered statistically significant.

Supporting Information

Figure S1 Effect of CR infection on NF-κB-p65 phosphorylation in Tlr4−/− mice. Representative photomicrographs of paraffin embedded sections prepared from the distal colons of uninfected normal (N) and days 9–19 post-CR infected Tlr4−/− mice and stained with H&E for gross morphology (left panel) and for p65 subunit phosphorylated at Ser-276 (p65527).
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Author Contributions
Conceived and designed the experiments: SA SU. Performed the experiments: CP LUMRJ BR IA. Analyzed the data: CP BR SU. Contributed reagents/materials/analysis tools: CP SU. Wrote the paper: CP SU.
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