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# *Campylobacter jejuni* Induces Colitis through Activation of Mammalian Target of Rapamycin Signaling

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# Abstract

**BACKGROUND & AIMS**—*Campylobacter jejuni* is the worldwide leading cause of bacterialinduced enteritis. The molecular and cellular events that lead to campylobacteriosis are poorly understood. We identify mammalian target of rapamycin (mTOR) as a signaling pathway that leads to *C jejuni*-induced intestinal inflammation.

**METHODS**—Germ-free (control) or conventionally-derived *interleukin* (*II*)10–/– mice that express enhanced green fluorescent protein under the control of NF- $\kappa$ B (*II10*–/–; NF- $\kappa$ B<sup>EGFP</sup> mice) were infected with *C jejuni* (109 CFU/mouse) for 12 days; their responses were determined using histologic, semi-quantitative reverse transcription PCR, fluorescence in situ hybridization, transmission electron microscopy, and tissue culture analyses. mTOR signaling was blocked by daily intraperitoneal injections of the pharmacologic inhibitor rapamycin (1.5 mg/kg). CD4<sup>+</sup> T cells were depleted by intraperitoneal injections of antibodies against CD4 (0.5 mg/mouse, every 3 days). Bacterial survival in splenocytes was measured using a gentamycin killing assay.

**RESULTS**—*C jejuni* induced intestinal inflammation, which correlated with activation of mTOR signaling and neutrophil infiltration. The inflamed intestines of these mice had increased levels of II-1 $\beta$ , Cxcl2, II-17a, and EGFP; *C jejuni* localized to colons and extra-intestinal tissues of infected *II10*—/–; *NF*- $\kappa B^{EGFP}$  mice, compared with controls. Rapamycin, administered before or after introduction of *C jejuni*, blocked *C jejuni*-induced intestinal inflammation and bacterial accumulation. LC3II processing and killing of *C jejuni* were increased in splenocytes incubated with rapamycin, compared with controls.

**CONCLUSIONS**—mTOR signaling mediates *C jejuni*-induced colitis in *Il10*–/– mice, independently of T-cell activation. Factors involved in mTOR signaling might be therapeutic targets for campylobacteriosis.

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## Keywords

crypt abscesses; imaging; mouse model; immune response; signal transduction

# INTRODUCTION

*Campylobacter jejuni* is a gram-negative category B priority food- and water-borne pathogen and the worldwide leading bacterial causative agent of enteritis.<sup>1</sup> The Centers for Disease Control and Prevention estimate that 2.4 million subjects are infected with *C. jejuni* resulting in 124 deaths every year in the United States.<sup>2</sup> ENREF 2 ENREF 2 Clinical symptoms of *C. jejuni* infection include abdominal cramps, watery to bloody diarrhea, fever and gastrointestinal inflammation.<sup>3</sup> At the cellular level, *C. jejuni* infected patients display infiltration of immune cells such as neutrophils, crypt abscesses and presence of fecal leukocytes.<sup>4</sup> Although the intestinal disease self-resolves within one week, a small portion of patients (1:1000) develop extra-intestinal sequelae such as Guillain-Barre Syndrome (GBS) and reactive arthritis.<sup>5</sup> Interestingly, *C. jejuni* exposure is also responsible for initiation and relapse of inflammatory bowel diseases (IBD)<sup>6</sup> and post-infectious irritable bowel syndrome<sup>7</sup>.

Despite the prevalence of *C. jejuni* induced illness and negative socio-economic impact, a paucity of information is available regarding the molecular and cellular events involved in campylobacteriosis. The major contributing factor to our poor understanding of campylobacteriosis is the lack of robust experimental models mimicking the various phases of acute human infection. Although some mammals including monkey,<sup>8</sup> ferret<sup>9</sup> and piglet<sup>10</sup> have provided valuable information regarding cellular events associated with campylobacteriosis, limited reagents and lack of genetic manipulation in these models have constrained the generation of deep mechanistic understanding. We recently established an acute model of campylobacteriosis using germ-free  $II10^{-/-}$  mice.<sup>11</sup> In this model, *C. jejuni* induces a rapid (5 days) and robust inflammatory (bloody diarrhea) response to the microorganism. However, the cellular and molecular details responsible for this host response remained undefined.

Mammalian target of rapamycin (mTOR) plays an important role in cell growth and proliferation.<sup>12</sup> The mTOR complex is composed of two entities: mTOR complex 1 (mTORC1) and mTOR complex 2 (mTORC2). mTORC1 is a downstream target of phosphoinositide 3-kinases (PI3K) and is sensitive to the pharmacological inhibitor rapamycin.<sup>13</sup> The PI3K/mTOR pathway has long been implicated in regulating adaptive immunity by targeting T cell proliferation and activation.<sup>14</sup> Recent evidence implicates PI3K/mTOR signaling as a key regulatory pathway of the innate immune host response. For example, rapamycin suppresses GM-CSF or IL-8 induced human neutrophil migration *in vitro*.<sup>15</sup> In addition, PI3K signaling mediates IL-17 expression in inflamed murine lungs.<sup>16</sup> Interestingly, IL-17 drives early neutrophil infiltration induced by i.p. *Escherichia coli* infection in mice.<sup>17</sup> Finally, mTOR signaling modulates autophagic responses involved in bacterial killing and clearance.<sup>18</sup>

In this study, we hypothesized that mTOR represents an important signaling event regulating host response to *C. jejuni* infection. Our findings demonstrated that mTOR signaling regulates *C. jejuni* induced expression of inflammatory mediators, neutrophil infiltration and bacterial clearance. These molecular and signaling events represent critical steps in *C. jejuni* induced intestinal inflammation and could define potential new therapeutic targets.

# MATERIALS AND METHODS

#### Mice

All animal protocols were approved by the Institutional Animal Care and Use Committee of the University of North Carolina at Chapel Hill. Germ-free 8-12 week old Il10<sup>-/-</sup>; NFκBEGFP (129/SvEv; C57BL/6 mixed background) mice were transferred from germ-free isolators and immediately gavaged with  $10^9$  C. *jejuni* cfu/mouse (strain 81–176<sup>19</sup>). Mice were then housed under Specific Pathogen-Free (SPF) conditions and sacrificed at 4 or 12 days. Germ-free  $Il10^{-/-}$ ; NF- $\kappa$ B<sup>EGFP</sup> mice were also gavaged with 10<sup>7</sup> Salmonella Typhimurium (S. Typhimurium strain CA32) cfu/mouse for 2 days or 10<sup>7</sup> Escherichia coli (E. coli strain NC 101) cfu/mouse for 12 days. Conventionally-derived Il10<sup>-/-</sup>; NFκB<sup>EGFP</sup> mice were put on an antibiotic cocktail (streptomycin 2 g/L, gentamycin 0.5 g/L, bacteriocin 1 g/L and ciprofloxacin 0.125 g/L) in water for 7 days and antibiotics were withdrawn 24 hr prior infection. The mice were then gavaged with 10<sup>9</sup> C. jejuni cfu/mouse for 14 days. To inhibit mTOR signaling, mice were injected daily i.p. with rapamycin (1.5 mg/kg) (Fisher Scientific). At the end of the experiment, mice were euthanized using CO<sub>2</sub> intoxication. Colon, mesenteric lymph nodes (MLN) and spleen were collected for further processing RNA, protein or C. jejuni culture on Campylobacter selective blood plates (Remel) for 48 h at 37 °C using the GasPak system (BD). Colons were fixed in 10% buffered formalin (Fisher Scientific) overnight, paraffin-embedded, sectioned, and stained with H&E for histological evaluation. Images were acquired using a DP71 camera and DP Controller 3.1.1.276 (Olympus). Intestinal inflammation was scored by evaluating the degree of lamina propria immune cell infiltration, goblet cell depletion, architectural distortion, as well as crypt hyperplasia, ulceration, and abscesses using a score from 0-4.11 Cecal and colonic inflammation induced by S. Typhimurium was evaluated using a score from 0–13 as described.<sup>20</sup>

#### CD4+ cell depletion

*C. jejuni* infected mice were injected with anti-CD4 antibody (BioXcell) (i.p. 0.5 mg/mouse, every 3 days) to deplete CD4+ cells. At the end of the experiment, mice were euthanized using CO<sub>2</sub> intoxication. Colons were resected and processed for H&E staining and colitis evaluation. Spleen and MLN were collected and processed into single cell suspensions after erythrocyte lysis.

#### Western blotting

Whole tissues were lysed in Laemmli buffer and 20  $\mu$ g of protein was separated by SDS-PAGE, transferred to nitrocellulose membranes and protein detected using enhanced chemiluminescence reaction (ECL) as described previously.<sup>11</sup> Primary antibodies used were phospho-p70S6K (T389) and p70S6K (Cell Signaling), EGFP (Sigma), LC3 I/II (MBL) and Actin (BP Biomedicals).

# Enhanced GFP (EGFP) macro- and micro-imaging

*Il10<sup>-/-</sup>*; NF-κB<sup>EGFP</sup> mice were sacrificed; the colon was dissected and immediately imaged using a charge-coupled device camera in a light-tight imaging box with a dual-filtered light source and emission filters specific for EGFP (LT-99D2 Illumatools; Lightools Research). For microimaging, colonic segments were cut open and fixed in 4% paraformaldehyde overnight at 4°C, and then permeabilized in 2% triton overnight at 4°C. The tissues were then immersed in FocusClear (CelExplorer labs) and then stained with DAPI. The tissues were put on slides with lumen side facing the lens. NF-κB derived EGPF were examined using Olympus FV1000 confocal multiphoton upright microscope system with Olympus

Fluoview 2.0 software. Acquired images were reconstructed into 3-D using Imaris (Bitplane).

## Fluorescence in situ hybridization (FISH)

Cy3-tagged 5'AGCTAACCACACCTTATACCG3'<sup>21</sup> was used to probe the presence of *C. jejuni*. Deparaffinized, formalin-fixed 5 µm thick sections were incubated for 15 minutes in lysozyme (300,000 Units/ml lysozyme; Sigma-Aldrich) buffer (25 mM Tris pH 7.5, 10 mM EDTA, 585 mM sucrose, and 0.3 mg/ml sodium taurocholate) at room temperature and hybridized overnight at 46 °C in hybridization chambers with the oligonucleotide probe (final concentration of 5 ng/µl in a solution of 30 percent formamide, 0.9 M sodium chloride, 20 mM Tris pH 7.5, and 0.01% sodium dodecyl sulfate). Tissue sections were washed for 20 minutes at 48 °C in washing buffer (0.9 M sodium chloride, 20 mM Tris pH 7.5) and once in distilled water for 10 seconds, dried at 46 °C, mounted with DAPI mount media, and imaged using a Zeiss LSM710 Spectral Confocal Laser Scanning Microscope system with ZEN 2008 software. Acquired images were analyzed using BioimageXD.<sup>22</sup> For whole tissue FISH, colons were fixed in 4% paraformaldehyde overnight at 46°C. The whole tissue was immersed in FocusClear, imaged on an Olympus FV1000 microscope and reconstructed into 3-D using Imaris.

#### Immunohistochemistry (IHC)

Deparaffinized colon sections were treated with 3% H<sub>2</sub>O<sub>2</sub> for 10 minutes to quench endogenous peroxidases. Antigen retrieval was performed using microwave heating for 10 min in 10 mM citrate buffer pH 6.0. Sections were blocked in 5% goat serum TBS-T for 1 hour. The slides were incubated overnight at 4°C with an anti-MPO antibody (1:400) (Thermo Scientific). Secondary biotinylated antibody was diluted at 1:1000 with the Vectastain ABC Elite Kit (Vector Laboratories). Visualization was performed using DAB chromogen (DAKO). Finally, sections were counterstained with hematoxylin for 5 seconds. Sections were imaged at the Olympus microscope using DP 71 camera and DP Controller 3.1.1.276 (Olympus).

#### Statistical analysis

Values are shown as mean  $\pm$  SEM as indicated. Differences between groups were evaluated with the nonparametric Mann-Whitney U test. Experiments were considered statistically significant if *P* values < 0.05. All calculations were performed using Prim 5.0 software.

# RESULTS

#### mTOR mediates C. jejuni-induced colitis

Although the role of mTOR has moved beyond that of a modulator of T cell function, the involvement of this multifunctional kinase in host responses to pathogenic bacteria has not been clearly explored. To investigate the impact of mTOR on *C. jejuni*-induced colitis *in vivo*, germ-free  $II10^{-/-}$ ; NF- $\kappa$ B<sup>EGFP</sup> mice were transferred to specific pathogen free (SPF) housing and immediately gavaged with *C. jejuni* (10<sup>9</sup> CFU/mouse) and then injected i.p. daily with either vehicle (5% DMSO PBS) or rapamycin (1.5 mg/kg body weight) for 12 days. Western blot analysis showed that rapamycin attenuated *C. jejuni* induced phosphorylation of p70S6 kinase (T389), a downstream target of mTOR (Fig. 1A). As previously reported, <sup>11</sup> $II10^{-/-}$ ; NF- $\kappa$ B<sup>EGFP</sup> mice infected with *C. jejuni* showed severe intestinal inflammation as seen by extensive immune cell infiltration, epithelial ulceration, goblet cell depletion and crypt hyperplasia and abscesses compared to uninfected mice (Fig. 1B and 1D). Interestingly, rapamycin blocked *C. jejuni*-induced intestinal inflammation in

 $II10^{-/-}$ ; NF- $\kappa$ B<sup>EGFP</sup> mice. Since germ free mice have an immature immune system, we next treated conventionally-derived  $II10^{-/-}$ ; NF- $\kappa$ B<sup>EGFP</sup> mice with antibiotics and then infected these mice with *C. jejuni* for 12 days. As observed in germ free mice, *C. jejuni* induced severe intestinal inflammation in conventionally-derived  $II10^{-/-}$ ; NF- $\kappa$ B<sup>EGFP</sup> mice, an effect significantly attenuated by rapamycin exposure (Supplemental Fig. S1a–b).

To determine the specificity of rapamycin on other colitogenic microorganisms, we infected germ free  $II10^{-/-}$ ; NF- $\kappa$ B<sup>EGFP</sup> mice with the enteric pathogens *Salmonella Typhimurium* (10<sup>7</sup> CFU/mouse) for 2 days or *Escherichia coli* NC 101 (10<sup>7</sup> CFU/mouse) for 12 days. Interestingly, *E. coli* infected  $II10^{-/-}$ ; NF- $\kappa$ B<sup>EGFP</sup> mice did not develop intestinal inflammation at this time point, suggesting a specific host response to *C.jejuni* infection (Supplemental Fig. S2a–b). However,  $II10^{-/-}$ ; NF- $\kappa$ B<sup>EGFP</sup> mice infected with *S. Typhimurium* developed severe cecal and colonic inflammation, which was strongly attenuated by rapamycin exposure (Supplemental Fig. S3). These results suggest that mTOR signaling mediates deleterious responses from different enteropathogenic microorganisms.

*C. jejuni* induced intestinal inflammation is evident as early as day 4 in  $II10^{-/-}$ ; NF- $\kappa$ B<sup>EGFP</sup> infected mice (Supplemental Fig. S4a–b).<sup>11</sup> To further test if rapamycin may be used as an intervention agent, we injected the inhibitor daily to *C. jejuni* infected (4 days)  $II10^{-/-}$ ; NF- $\kappa$ B<sup>EGFP</sup> mice and collected intestinal tissues at 12 days. Remarkably, rapamycin treatment strongly reversed *C. jejuni* induced intestinal inflammation and bloody diarrhea (Fig. 1C and 1E, Supplemental Fig. S5a–b).

Since rapamycin has immunomodulatory effects on T cells, we depleted CD4+ T cells using an anti-CD4 antibody (0.5 mg/mouse/every three days) to address the function of these cells in the acute phase of campylobacteriosis. Notably, *C. jejuni* induced colitis was only slightly inhibited (~20%) in CD4+ T cell-depleted mice compared to control (Fig. 2A–B). Complete depletion of CD4 cells was observed in the spleen and MLN cellular compartment of antibody-treated mice (Fig. 2C–D). These results suggest that the early phase of *C. jejuni*induced colitis is mostly mediated by an innate immune response and that rapamycin likely targets activities of innate immune cells.

#### mTOR regulates C. jejuni induced NF-kB activity and proinflammatory gene expression

We previously showed that NF- $\kappa$ B is activated in *C. jejuni* infected *ll10<sup>-/-</sup>*; NF- $\kappa$ B<sup>EGFP</sup> mice.<sup>11</sup> To evaluate the impact of mTOR signaling on NF- $\kappa$ B activity, EGFP expression in the colon of *ll10<sup>-/-</sup>*; NF- $\kappa$ B<sup>EGFP</sup> mice was visualized using a CCD camera macroimaging system. *ll10<sup>-/-</sup>*; NF- $\kappa$ B<sup>EGFP</sup> mice infected with *C. jejuni* displayed enhanced colonic NF- $\kappa$ B<sup>EGFP</sup> expression with strongly positive lymphoid aggregates (arrow heads) compared to uninfected mice (Fig. 3A). Interestingly, the colon of rapamycin-treated, *C. jejuni*-infected *ll10<sup>-/-</sup>*; NF- $\kappa$ B<sup>EGFP</sup> mice displayed reduced EGFP expression compared to untreated, infected-mice. Consistent with these findings, western blot analysis of colonic lysates demonstrated reduced EGFP expression in rapamycin-treated, *C. jejuni*-infected mice (Fig. 3B).

To further evaluate the distribution of NF- $\kappa$ B<sup>EGFP</sup> positive cells within the colon, whole colonic tissue was visualized using confocal microscopy coupled with 3-D image reconstruction. In accordance with the macroimaging and Western blot data, *C. jejuni*-infected *II10<sup>-/-</sup>*; NF- $\kappa$ B<sup>EGFP</sup> mice showed numerous EGFP positive cells in the colonic lamina propria compared to uninfected mice (Fig. 3C and supplementary Fig3video1.mov, Fig3video2.mov, Fig3video3.mov and Fig3video4.mov). Strikingly, colonic EGFP tissue expression was strongly reduced in rapamycin-treated, *C. jejuni*-infected *II10<sup>-/-</sup>*; NF- $\kappa$ B<sup>EGFP</sup> mice compared to untreated mice.

We next examined the impact of rapamycin on expression of various NF- $\kappa$ B dependent proinflammatory mediators involved in bacterial host responses. *C. jejuni* infection strongly induced *II-1* $\beta$ , *Cxcl2* and *II-17a* mRNA accumulation in colonic tissue of *II10<sup>-/-</sup>*; NF- $\kappa$ B<sup>EGFP</sup> mice, effects attenuated by 79, 75, and 92%, respectively in rapamycin-treated, *C. jejuni*-infected mice (Fig. 4A). Consistent with these findings, secreted IL-1 $\beta$  and IL-17 from supernatant of cultured colon explants was reduced by 59% and 94% respectively while production of these cytokines was attenuated by 41% and 82% in MLN of rapamycintreated, *C. jejuni*-infected mice compared to control mice (Fig. 4B). Interestingly, *C. jejuni*induced colonic *II-12p40* and *Tnf* $\alpha$  mRNA accumulation were not significantly inhibited by rapamycin treatment, suggesting that inflammatory mediators are selectively affected by mTOR signaling (Supplemental Fig. S6).

## Neutrophils participate in C. jejuni-induced colitis

Neutrophil infiltration is a hallmark of acute infection and represents an important feature of human campylobacteriosis.<sup>4</sup> Of note, rapamycin strongly reduced *C. jejuni* induced expression of the neutrophil marker myeloperoxidase (MPO) in colonic tissue of infected  $II10^{-/-}$ ; NF- $\kappa$ B<sup>EGFP</sup> mice (Fig. 5A). Moreover, the ratio of Gr-1<sup>+</sup>:CD45<sup>+</sup> cells significantly decreased (41%) in the blood of rapacymin-treated, *C. jejuni* infected mice compared to untreated, infected mice (Fig. 5B). Finally, transmission electron microscope (TEM) analysis showed neutrophil accumulation in colonic crypt, which associates with microvilli/glycocalyx destruction (Fig. 5C). Notably, these pathological features were not observed in rapamycin-treated mice (Fig. 5C, right panel).

#### mTOR mediates C. jejuni invasion

We next investigated the impact of mTOR signaling on intestinal and extra-intestinal *C. jejuni* tissue distribution. Following infection and treatment with rapamycin, *C. jejuni* DNA was visualized in the colon of  $II10^{-/-}$ ; NF- $\kappa$ B<sup>EGFP</sup> mice using fluorescence *in situ* hybridization (FISH) and confocal microscopy imaging. Interestingly, while *C. jejuni* was detected deeply in the inflamed crypts and in the lamina propria section of the intestine of untreated mice, *C. jejuni* DNA was barely detectable in rapamycin-treated mice (Fig. 6A).

To gain a better perspective of *C. jejuni* invasion, FISH was performed using fresh-fixed colonic tissues sections, and then imaged with confocal microscopy coupled with 3-D visualization. Interestingly, while *C. jejuni* extensively invaded the colon of untreated mice, its presence is strongly reduced in rapamycin-treated mice (Fig. 6B and supplementary Fig6video1.mov and Fig6video2.mov). To further characterize *C. jejuni* invasion, colonic tissue was also imaged using TEM. The presence of spirally shaped *C. jejuni* (arrow) lacking its outer membrane was detected in epithelial cells (Fig. 6C left panel). Also the absence of a vesicular membrane around the invading bacteria suggests that phagosome formation in epithelial cells is not a host response to the bacterium. Consistent with FISH results, intracellular *C. jejuni* were virtually absent and microvilli were intact in epithelial cells of rapamycin-treated mice (Fig. 6C right panel).

Reduced *C. jejuni* in the colon of rapamycin-treated mice suggests that the bacteria were eliminated or were able to evade/translocate to extra-intestinal tissues. To resolve this issue, we aseptically collected samples from the stool, colon, spleen and MLN and enumerated *C. jejuni* on Remel *Campylobacter* selective plates. Consistent with the FISH and TEM results, rapamycin treatment reduced *C. jejuni* colonic tissue invasion by 90% compared to *C. jejuni*-infected, untreated mice (Fig. 6D). Furthermore, rapamycin treatment strongly reduced *C. jejuni* presence in the MLN and spleen of infected mice compared to untreated, infected mice. Altogether, these findings identified mTOR signaling and neutrophil infiltration as important events leading to *C. jejuni* invasion and pathogenesis.

The above findings suggest that rapamycin favors bacterial clearance, which lead to decreased innate response and intestinal inflammation. To directly test the impact of rapamycin on bacterial survival, we infected splenocytes isolated from  $Il10^{-/-}$  mice with *C. jejuni* and performed a gentamycin killing assay. Interestingly, rapamycin enhanced *C. jejuni* killing in splenocytes at 4hr compared to untreated cell (Fig. 7A). Western blot analysis showed that rapamycin attenuated *C. jejuni* induced phosphorylation of p70S6 kinase (T389) and enhanced LC3 II conversion, a marker of autophagy (Fig.7B). Interestingly, *C. jejuni* exposed to rapamycin as high as 1 mM did not alter the growth/ viability of the bacterium. Altogether, these findings showed that increased mTOR signaling following *C. jejuni* infection mediates proinflammatory response, likely by preventing host-mediated bacterial killing.

# DISCUSSION

The fundamental molecular host response to C. jejuni infection remains virtually unknown. In this study, we present clear evidences that C. jejuni -mediated intestinal inflammation is caused by activation of the host mTOR signaling pathway and neutrophil infiltration. Comparable intestinal inflammation between CD4+ cells depleted and untreated mice indicates that *C. jejuni* induced colitis is mostly driven by innate immune cells at day 12. Although CD4+ T cells are not involved in the development of intestinal inflammation by day 12, adaptive host response is an important hallmark of campylobacteriosis.<sup>23</sup> Interestingly and in contrast to C. jejuni, the adherent invasive E. coli NC101 failed to induce colitis after 12 days, whereas severe colitis developed after 10-20 weeks of colonization with the same bacterium.<sup>24</sup> In addition, colitis was observed as early as two days in S. Typhimurium infected  $II10^{-/-}$  mice. These findings suggest a differential ability of bacteria to trigger intestinal inflammation in  $Il10^{-/-}$  mice. Using this robust and tractable animal model, we demonstrate that C. jejuni induced mTOR signaling leads to increased NF-κB activity and induction of NF-κB-dependent genes (cytokines/chemokines). FISH assay and electron microscopy analysis showed that C. jejuni rapidly invades the intestinal mucosal layer and triggers a host response involving neutrophil recruitment and formation of crypt ulcers. Disruption of the crypt architecture by massive neutrophil influx likely compromises the intestinal barrier integrity, which further favors bacterial uptake/invasion and dissemination to extra-intestinal tissues as seen by enhanced bacterial counts in the spleen and MLN. mTOR blockade with a pharmacological inhibitor strongly attenuated campylobacteriosis in  $Il10^{-/-}$  mice. Moreover, rapamycin was able to treat established C. *jejuni*-induced intestinal inflammation.

A close examination of human campylobacteriosis suggests a preponderant role of neutrophils in the course of the pathology. Histological assessment showed the presence of numerous crypt abscesses and neutrophil infiltration in the intestinal mucosa of infected patients.<sup>4</sup> However, up until now the functional importance of neutrophil infiltration in C. jejuni-induced pathogenesis has not been investigated. Although the main biological function of neutrophils is to ingest and eliminate invading microorganisms, excessive infiltration of these innate immune cells and release of various degradative enzymes and oxidative products cause extensive collateral tissue damage to the host. For example, neutrophil accumulation causes elevated IL-1 $\beta$  and CXCL2 production and subsequent joint inflammation, an effect attenuated with anti-IL1ß antibody treatment.<sup>25</sup> Interestingly, TEM showed that the microvilli and associated glycocalyx of epithelial cells in neutrophilinfiltrated crypts are virtually ablated in C. jejuni-infected mice.-The molecular events leading to neutrophil infiltration in the intestine of C. *jejuni*-infected mice is not entirely clear. Induction of Il-17a and Cxcl2 genes is important for neutrophil expansion and recruitment.<sup>26, 27</sup> Interestingly, blocking mTOR with rapamycin decreased colonic *Il-17a* and Cxcl2 expression, which correlated with reduction of neutrophils infiltration and crypt

abscesses and with disease improvement. Although IL-17 is a signature cytokine of the Th17 cells, our CD4 depletion experiment suggests that adaptive CD4<sup>+</sup> T cells play a minor role in campylobacteriosis. IL-17 is produced by a wide array of innate immune cells including  $\gamma\delta T$  cells, natural killer  $\gamma\delta T$  cells and lymphoid tissue inducer cells.<sup>28</sup> Since  $\gamma\delta T$  cells are highly abundant in the intestinal mucosa, it is tempting to speculate that these cells are the source of IL-17 in our model. Interestingly, in an *E. coli* infectious model, neutrophil recruitment to the peritoneal cavity is dependent on innate  $\gamma\delta T$  cell-mediated IL-17 secretion.<sup>17</sup> Similarly, *Klebsiella pneumonia*-induced lung inflammation is attenuated in *Il-17r<sup>-/-</sup>* mice, which correlated with neutrophil recruitment and G-SCF and CXCL2 expression.<sup>27</sup> Further studies would be needed to define the role of IL-17 in campylobacteriosis.

Another important function of mTOR signaling is the regulation of autophagy. Our findings indicate that C. jejuni infiltration in the colon, MLN and spleen is dramatically reduced in rapamycin-treated mice, suggesting increased bacterial killing. Using primary splenocytes, we found that rapamycin enhanced C. jejuni killing, which correlated with enhanced LC3II conversion. These findings suggest that C. jejuni induces mTOR signaling as a mean to avoid host-mediated killing, possibly by prevent adequate autophagy response. Further studies would be needed to definitively establish the role of autophagy in campylobacteriosis. Interestingly, S. Typhimurium induced colitis was also inhibited by rapamycin exposure, suggesting that mTOR may be the target of other enteropathogenic microorganisms. mTOR has also been recently identified as a negative regulator of LPSinduced NF-KB signaling in monocytes, macrophage, and primary DC.<sup>29</sup> Therefore, blocking mTOR signaling could enhance NF-κB signaling and production of inflammatory mediators involved in bacterial killing. However, our data do not support this hypothesis as EGFP expression (NF-KB activity) and some NF-KB-dependent inflammatory mediators are reduced in rapamycin-treated mice. While we observed decreased activation of NF-kB (EGFP expression) and inflammatory gene expression in rapamycin-treated mice, these observations are likely a secondary effect of the drug since direct NF-kB inhibitors failed to prevent campylobacteriosis.<sup>11</sup>

Although the majority of *C. jejuni* infected patients recover within 2 to 10 days without any specific treatment, a significant proportion develop severe clinical symptoms (fever, bloody diarrhea and abdominal cramps) which requires antibiotic treatment to shorten symptom durance.<sup>1</sup> Recently, there are increasing concerns over *C. jejuni* resistance to antibiotics<sup>30</sup> and post-infectious sequelae such as GBS, IBD and IBS.<sup>5–7</sup> Immune based prevention or intervention has attracted much attention as an alternative to antibiotic treatment.<sup>31</sup> Based on our novel findings, specific mTOR inhibitors (such as Rapamune, Torisel, Afinitor and Zortress) already approved for human use could represent a potential alternative to antibiotic-based therapy to treat campylobacteriosis. Further work would be needed before mTOR is validated as a therapeutic target for campylobacteriosis.

In summary, this report defines the critical role of mTOR in mediating the pro-inflammatory effect of *C. jejuni* infection. We report that neutrophil infiltration plays an active role in pathogenesis of *C. jejuni* infection and modulation of the cellular/molecular events leading to this process could represent a new therapeutic arsenal to control campylobacteriosis.

# Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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# Abbreviations

mTOR	mammalian target of rapamycin
MLN	mesenteric lymph nodes
IHC	immunohistochemistry
FISH	Fluorescence in situ hybridization
TEM	transmission electron microscope
ELISA	Enzyme-linked immunosorbent assay

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**Figure 1. Rapamycin prevents and treats** *C. jejuni* induced colitis in  $II10^{-/-}$ ; NF- $\kappa$ B<sup>EGFP</sup> mice Cohorts of 5–14 germ-free  $Il10^{-/-}$ ; NF- $\kappa$ B<sup>EGFP</sup> mice were transferred to SPF conditions and immediately gavaged with 10<sup>9</sup> *C. jejuni*/mouse and then injected i.p. with vehicle control (Ctl, 5% DMSO), rapamycin (Rapa, 1.5 mg/kg body weight) for 12 days. For rapamycin treatment (Rapa trt), *C. jejuni*-infected mice (4 days) were injected i.p. with rapamycin to day 12. Colon was resected, fixed in formaldehyde and sections were stained with H&E or protein extracts for Western blot analysis. (A) Western blot for total and phosphorylated colonic p70S6. (B) Histological intestinal damage scores of rapamycin treatment. (D) H&E representation of the different experimental groups with crypt abscesses (arrow heads). (E) H&E sections of rapamycin treatment mice. All graphs depict mean  $\pm$  SE. \*\*, P < 0.01, \*\*\*, P < 0.001. Scale bar is 200 µm. Results are representative of 4 independent experiments.



Figure 2. *C. jejuni* induced intestinal inflammation is independent of CD4 T cells activation Three cohorts of 4–5 germ-free *Il10<sup>-/-</sup>*; NF- $\kappa$ B<sup>EGFP</sup> mice were transferred to SPF conditions and immediately gavaged with 10<sup>9</sup> *C. jejuni*/mouse and then injected i.p. with vehicle control (Ctl, PBS) or anti-CD4 antibody (anti-CD4, 0.5 mg/mouse/every 3 days). (A) H&E representation of the different experimental groups. (B) Histological score of intestinal inflammation. (C) Representative flow cytometry analysis of CD4+ cell population change in the spleen. (D) Percentage flow cytometry results of CD4+ cell in the spleen and MLN. All graphs depict mean ± SE. \*\*\*, P < 0.001. Scale bar is 200 µm. Results are representative of 3 independent experiments.



Figure 3. Rapamycin treatment reduces *C. jejuni* induced colonic EGFP expression in *Il10*<sup>-/-</sup>; NF-κB<sup>EGFP</sup> mice

Four cohorts of 5–14 germ-free  $ll10^{-/-}$ ; NF- $\kappa$ B<sup>EGFP</sup> mice were transferred to SPF conditions and infected with *C. jejuni*/mice as described in Fig.1. Mice were injected i.p. with vehicle (5% DMSO) or rapamycin (1.5 mg/kg body weight) for 12 days. (A) EGFP expression in the mouse colon was visualized using a CCD camera macroimaging system. (B) EGFP levels were analyzed by Western blot and (C) confocal microscopy visualization of EGFP positive cells images in 3-D using Imaris. Results are representative of 3 independent experiments.



**Figure 4. Rapamycin attenuates** *C. jejuni* induced expression of proinflammatory mediators Four cohorts of 5–14 germ-free  $II10^{-/-}$ ; NF- $\kappa$ B<sup>EGFP</sup> mice were treated as indicated in Fig.1. Colons were resected and RNA extracted using Trizol. (A)  $II1\beta$ , *Cxcl2* and *Il-17a* mRNA accumulation was quantified using an ABI 7900HT Fast Real-Time PCR System and specific primers and data were normalized to *Gapdh*. (B) Colonic tissues and mesenteric lymph nodes were cultured in RPMI 1640 medium supplemented with 3% FBS and 1% antibiotics for 18 hrs. Supernatants were collected and IL-1 $\beta$  and IL-17 secretion measured by ELISA. Data represent means ± SE \*, P < 0.05 \*\*, P < 0.01. Results are representative of 3 independent experiments.





Cohorts of 5–7 germ-free  $II10^{-/-}$ ; NF-KB<sup>EGFP</sup> mice were transferred to SPF conditions and infected/treated as indicated in Fig.1. (A) IHC representation of MPO positive colonic cells (brown dots) in Control (Ctl), *C. jejuni* infected, rapamycin (Rapa) treated mice. Lower panels (scale bar 20 µm) are magnified images of area shown in the upper panels (scale bar 200 µm). (B) Flow cytometry analysis for CD45<sup>+</sup> and Gr-1<sup>+</sup> cells in the peripheral blood of infected and rapamycin-treated mice. (C) TEM images of colonic tissues from *C. jejuni* infected with/without rapamycin treatment. Left panel shows accumulated neutrophils (arrow heads) in a crypt with microvilli erosion (arrows). Right panel shows absence of neutrophils with intact microvilli (arrows). Inserts in the lower right corners are magnified images of framed area. Scale bar is 2 µm. Data represent means ± SE \*\*, P < 0.01. Results are representative of 3 independent experiments.







#### Figure 7. Rapamycin promotes *C. jejuni* eradication and LC3II generation in splenocytes

Splenocyte was isolated from  $II10^{-/-}$  mice and infected with *C. jejuni* (MOI 50) and cultured with rapamycin (100 nM). A gentamycin killing assay at 0 and 4 hr and WB assay at 0.5, 1.5 and 4 hr were performed. (A) Percentage of *C. jejuni* survival in splenocytes. (B) Western blot of colonic LC3 I/II and phosphorylated p70S6 in infected splenocytes.