



Published in final edited form as:

Auton Neurosci. 2018 September ; 213: 51–59. doi:10.1016/j.autneu.2018.05.009.

S. Typhimurium challenge in juvenile pigs modulates the expression and localization of enteric cholinergic proteins and correlates with mucosal injury and inflammation

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Abstract

The cholinergic system plays a central role in regulating critical gastrointestinal functions, including motility, secretion, barrier and immune function. In rodent models of acute, non-infectious gastrointestinal injury, the cholinergic system functions to inhibit inflammation; however, during inflammation local expression and regulation of the cholinergic system is not well known, particularly during infectious enteritis. The objective of this study was to determine the intrinsic expression of the enteric cholinergic system in pig ileum following an acute challenge with *Salmonella enterica* serovar Typhimurium DT104 (*S. Typhimurium*). At 2 d post-challenge, a three-fold reduction in ileal acetylcholine (ACh) levels was observed in challenged animals, compared with controls. Ileal acetylcholinesterase (AChE) activity was decreased (by four-fold) while choline acetyltransferase (ChAT) expression was increased in both the ileum and mesenteric lymph nodes. Elevated ChAT found to localize preferentially to mucosa overlying lymphoid follicles of the Peyer's patch in challenged pigs, with more intense labeling for ChAT in *S. Typhimurium* challenged pigs compared to controls. Ileal mRNA gene expression of muscarinic receptor 1 and 3 was also increased in challenged pigs, while muscarinic receptor 2 and the nicotinic receptor alpha 7 subunit gene expression were unaffected. A positive correlation was observed between ChAT protein expression in the ileum, rectal temperature, and histopathological severity in challenged animals. These data show that inflammation from *S. Typhimurium* challenge alters enteric cholinergic expression by down-regulating acetylcholine concentration and acetylcholine degrading enzymes while increasing acetylcholine synthesis proteins and receptors. Given the known anti-inflammatory role of the cholinergic system, the divergent expression of

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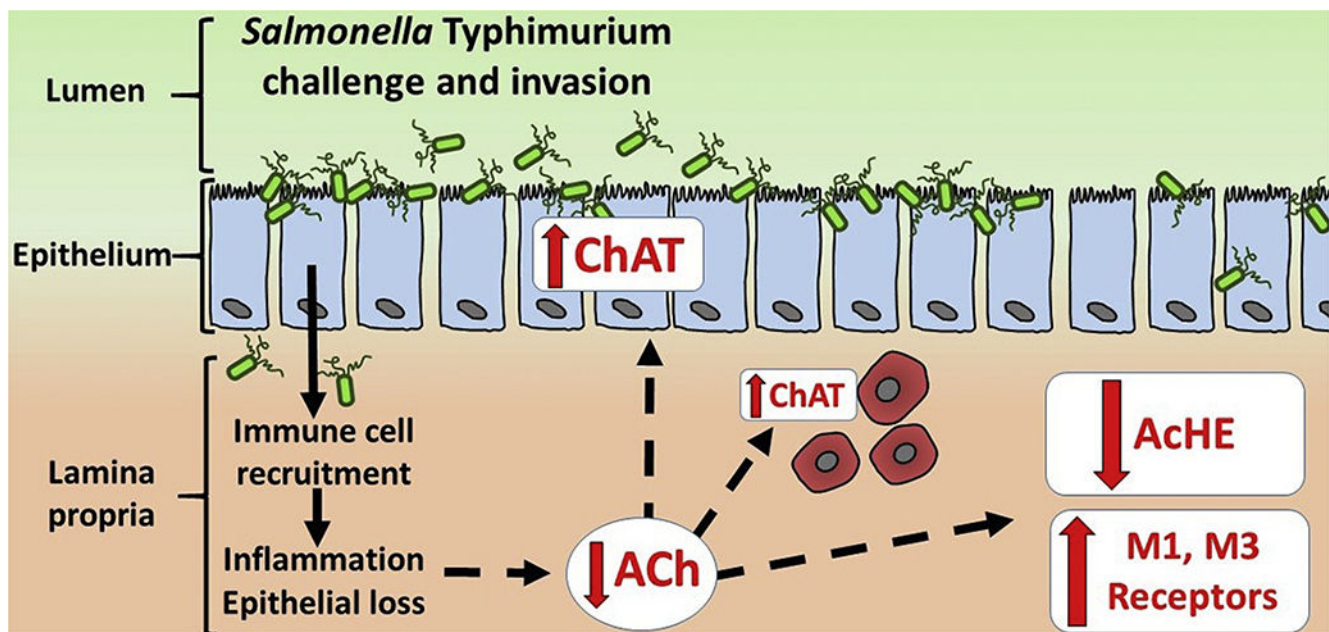
Conflicts of Interest: None.

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cholinergic genes may represent an attempt to limit tissue damage by preserving cholinergic signaling in the face of low ligand availability.

Graphical Abstract. Alterations in the enteric cholinergic system induced by *Salmonella* Typhimurium challenge in pigs.

Salmonella Typhimurium challenge and invasion triggers immune cell recruitment, inflammation and mucosal injury including epithelial cell loss. Subsequent reduction in acetylcholine (ACh) concentrations in the mucosa may induce a compensatory response to preserve and (or) amplify cholinergic signaling by (1) upregulating ACh synthesis via choline acetyltransferase (ChAT) in epithelial cells, lamina propria monocytes and enteric neurons (not shown) (2) down-regulating ACh degradation via suppression of acetylcholinesterase (AChE), and (3) enhancing of ACh signaling via increased expression of muscarinic M1 and M3 cholinergic receptors. Dotted lines indicate hypothesized mechanisms by which mucosal damage and inflammation may drive altered cholinergic expression.



Keywords

Cholinergic; Mucosal Inflammation; *Salmonella enterica* Typhimurium; porcine intestine

1. Introduction

To survive infectious challenges, the host must balance pathogen inducing immunity and inflammation with responses that limit tissue damage. Therapeutic interest focuses on pathways that simultaneously limit tissue damage without comprising sterilizing immunity (Schneider and Ayres 2008; Soares et al. 2014). Due to the intimate juxtaposition of microbes and host tissue, the gastrointestinal tract is an area requiring precise balance

between immunity and host tissue survival, and the cholinergic system is considered to be an important factor mediating this balancing act during enteritis (Gabanyi et al. 2016).

Cholinergic signaling via the vagus nerve is a well-established mediator of inflammation in the gastrointestinal tract (Matteoli and Boeckxstaens 2013). Several groups have shown that cholinergic function, mostly through action of the vagus nerve and nicotinic receptors, reduces gastrointestinal inflammation and prevents mucosal tissue damage in rodent models of chemical colitis (Ghia et al. 2006; Ghia et al. 2008; Mazelin et al. 1998; O'Mahony et al. 2009), endotoxemia (de Haan et al. 2013), and postoperative ileus (Matteoli et al. 2014; The et al. 2011). Further anti-inflammatory influence of the cholinergic vagal system is known by its role in promoting oral tolerance to foreign antigens (Di Giovangiulio et al. 2016).

The functional role of the cholinergic system in infectious enteritis is less well described; however, initial reports suggest that in contrast to the chemical colitis and postoperative ileus models, cholinergic signaling may enhance immunity by promoting inflammation. For example, pretreating mice with an acetylcholinesterase inhibitor prior to oral *Salmonella* infection increased serum inflammatory cytokine production, bacterial clearance, and host survival (Fernandez-Cabezudo et al. 2010). Another report demonstrated that cholinergic muscarinic receptors enhance T-cell pro-inflammatory activity and contribute to rapid convalescence and generation of adaptive immunity against both bacterial and parasitic infection (Darby et al. 2015). However, certain bacterial and parasitic, infectious enteritis models were also shown to result in suppressed cholinergic enteric nerve activity and attenuated release of Ach (Bercik et al. 2002; Collins et al. 1989; Galeazzi et al. 2000). Together, previous studies in rodents indicate that the cholinergic system has significant, yet divergent actions during mucosal inflammation which may depend upon the inciting stimuli.

The cholinergic system has several components that regulate the synthesis, degradation and signaling of Ach. Acetylcholine (ACh), the primary endogenous ligand inducing cholinergic signaling can be synthesized in both neuronal and non-neuronal cells. Choline acetyltransferase (ChAT), is the primary enzyme that generates acetylcholine, and it can be found in epithelia (Gareau et al. 2007), immune (Dhawan et al. 2016; Kawashima et al. 2007; Reardon et al. 2013), and neuronal cells (Vieira et al. 2017; Winston et al. 2013). Once acetylcholine is liberated into the extracellular space, it acts on two classes of cholinergic receptors, known as muscarinic and nicotinic receptors. Muscarinic receptors are G-coupled protein receptors that have up to 5 different sub-classes (Kruse et al. 2014). Nicotinic receptors are pentameric, ionotropic receptors that are formed by heterogeneous or homogenous assembly of one of several different nicotinic subunits (Albuquerque et al. 2009). Both types of receptors can be found on several different cells, mediating vastly different homeostatic functions. Finally, cholinergic signaling is terminated primarily by acetylcholinesterase (AChE), an enzyme that breaks down ACh at high efficiency and is expressed in many different cell types. While the functions of each component of the cholinergic system is well-established, how each component is dynamically during infectious challenge remain poorly defined. Considering the significant contribution of acetylcholine to gut homeostasis, understanding the expression of the cholinergic system during inflammatory challenges will provide a foundational understanding for future research and therapies. Therefore, In the present study characterized the changes in

expression of the enteric mucosal cholinergic system components in pigs acutely challenged with *S. Typhimurium* and how these changes are correlated with tissue damage and mucosal inflammatory cytokine production.

Materials and Methods

2.1. Animals and experimental design

Data was generated from tissues which were collected in a previously reported experiment (Boyer et al. 2015), which was under an approved Institutional Animal care and Use Committee at North Carolina State University (protocol no. 12–051-A). As reported previously, animals used were Yorkshire-Large White piglets weaned at 16–17 days of age and housed at 8 pigs per pen with *ad libitum* access to water and feed. Sex was distributed equally across weaning groups. At 34 days post weaning, all piglets were transferred to isolation rooms and housed by treatment groups with continued *ab libitum* access to food and water. *S. Typhimurium* challenged pigs were orally inoculated with 3×10^9 colony forming units in 4mls of culture media; while uninfected controls were feed 4mls of sterile culture media. Challenged pigs were housed in separate rooms from the unchallenged controls; however the housing and environmental conditions were identical between rooms. *S. Typhimurium* DT104 strain culture was grown overnight at 37°C in Luria broth agar and added to 0.7% sterile saline for form final concentrations of 7.5×10^8 colony forming unites per mL (Boyer et al. 2015). Animals were euthanized 2 days post pathogen challenge, and collection of ileum mucosal scrapes and lymph nodes were performed and stored as previously reported (Boyer et al. 2015). The post-challenge time point (48 h) was selected to coincide with the peak acute phase responses (rectal temperature, feed intake reduction, plasma cortisol, etc.) to oral *S. Typhimurium* challenge in pigs as reported previously (Balaji et al. 2000).

2.2 Ileum and mesenteric lymph node protein Isolation

For SDS-PAGE and Western blot, 0.5cm³ pieces of ileal mucosa scrapes and mesenteric lymph node were collected over dry ice and homogenized in RIPA buffer (Thermo Scientific, #89900) in the presence of 1x protease inhibitor cocktail (Sigma Aldrich, #P8340) and 1x Halt Phosphatase Inhibitor (Thermo Scientific, #78420). Samples were spun at 13,300rpm at 4°C for 15min. Supernatant was collected, aliquoted and frozen at –70°C. Protein concentration was determine with Pierce BCA kit (Thermo Scientific, #23225), and samples were diluted to working a concentration of 2µg/µl.

For TNF and IL-8 ELISA and myeloperoxidase assay, samples were isolated as previously reported (Boyer et al. 2015).

2.3 SDS PAGE | Western Blot

Ileal mucosa and mesenteric lymph node protein samples were diluted to 1µg/µL in Laemmli Buffer (Bio-Rad, #161–0737) + 5% 2-mercaptoethanol and heat denatured at 70°C for 10 minutes. 10 µg of protein sample was run on a TGX-Stain Free gel (Bio-Rad #5678095). Protocol for electrophoresis, wet to wet transfer, and stain free, lane total protein quantification was performed as published in Criterion™ Precast Gels: Instruction Manual

and Application Guide and Western Blot Normalization Using Image Lab™ Software (Bio-Rad). The PVDF membrane was blocked in 5% BSA at RT for 1hr prior to incubation with monoclonal 1.B3.9B3 anti-porcine ChAT antibody (Millipore Sigma #MAB5270) at a concentration of 1:1000 in 1xTBS + 5% BSA + 0.1% Tween-20 overnight at 4°C. The following morning, the blot was washed and an HRP linked anti-mouse antibody in 1xTBS + 5% BSA + 0.1% Tween-20 (Cell Signaling, #7076) at (1:1000) was incubated with the membrane for 1hr at RT. Chemiluminescence was performed using Clarity ECL (Bio-Rad, #1705060). Densitometry was performed utilizing Bio-Rad Image Lab™ Software v5.2.1 and band density was normalized to lane total protein per Western Blot Normalization Using Image Lab™ Software (Bio-Rad) protocols. Total protein image for corresponding lanes in Figure 1 are in Supplemental Figure 1.

2.4 Acetylcholine quantification and acetylcholinesterase activity

Acetylcholine and acetylcholinesterase activity was performed on protein isolated from ileum mucosal scrapes using Amplex™ Acetylcholine/Acetylcholinesterase Assay Kit (ThermoFisher Scientific cat#A12217) as per manufacturer's instructions.

2.5 Gene Expression Analysis

Total RNA samples were isolated from frozen intestinal mucosal scrapings using the Qiagen RNeasy Mini kit. First-strand cDNA was synthesized from 4 µg RNA using Thermo Scientific Maxima First Strand cDNA Synthesis Kit for RT-qPCR with dsDNase (Thermo Scientific, K1641) according to the manufacturer's instructions. Semi-quantitative, real-time PCR was used to determine the relative quantities of transcripts of the genes of interest. Beta-actin (*ACTB*) was selected and validated as suitable internal reference genes. The relative gene expressions of cholinergic receptor muscarinic 1 (*CHRM1*), 2 (*CHRM2*), and 3 (*CHRM3*), cholinergic receptor nicotinic alpha 7 subunit (*CHRNA7*) were determined. Primer sequences for all genes are provided in Supplemental Table 1. All PCR reactions were subjected to a melt curve analysis to validate the absence of nonspecific products. The data are presented as 2^{-CT} in gene expression relative to control group, normalized to the *ACTB* before statistical analysis.

2.6 Correlation Analysis

Correlations were performed between mucosal acetylcholine concentrations, acetylcholinesterase activity, and the ChAT band identified on Western blot and rectal temperature, ileum histopathological scores, or ileum mucosal cytokine levels reported in a previous publication (Boyer et al. 2015). For methods on histopathology and mucosal cytokine analysis, please refer to our previous manuscript. (Boyer et al. 2015) Comparisons between ChAT and cytokines of interest was performed per each animal, and two tailed Pearson correlations were run on each group to identify any positive or negative association between ChAT expression and cytokine protein.

2.7 Immunohistochemistry and Image analysis

Sections of ileum were fixed in 10% neutral buffered formalin and paraffin embedded. Sections were prepped and immunohistochemically labeled for ChAT with B3.9B3 anti-

porcine ChAT antibody (Millipore Sigma #MAB5270) at 1:100. Detection of ChAT in section was performed by using secondary anti-mouse-on-Farma HRP polymer for 30min at RT and treatment with Romulin AEC. Slides were counter stained with hematoxylin at 1:10. All sample preparation and labeling was performed by Michigan State University's Investigative Histopathology Laboratory. Total mucosal area of ChAT positive labelling and integrated density of ChAT labelling were determined to generally assess the number of cells positive to ChAT and the intensity with which ChAT was expressed, respectively. Using ImageJ (U.S. NIH, Bethesda, MD, USA), the total area and integrated density of ChAT expression was determined by using the threshold function on the blue stack for each RGB image. Using the threshold function, we were able to assess the area and integrated density of the ChAT labeling alone, independent of the hematoxylin counter stain. The threshold was set from 0–61 and was consistently used on each image assess. The percentage area of ChAT positive staining was determined by dividing the area of ChAT positive labeling by the total area of the mucosa in the field. A total of 10 different tissue sections per slide were analyzed from each slide/animal and then average to derive the mean for individual animal.

2.8 Statistics

Two-tailed student's t-test were performed on most data comparing unchallenged controls to *S. Typhimurium* challenged animals. Unless reported otherwise, all values reported are means and standard error of means. Two tailed student t-tests and Pearson correlations and figures were generated with GraphPad Prism 6, v6.04 (GraphPad Prism Software).

3. Results

3.1 *Summary of S. Typhimurium clinical and histopathological findings as previously reported*

In Table 1, we summarize clinical and ileal histopathological features between uninfected controls and *S. Typhimurium* challenged animals as previously reported (Boyer et al. 2015). *S. Typhimurium* challenge induced significant diarrhea and pyrexia. Histopathology scores demonstrate that *S. Typhimurium* induced intestinal mucosal injury with moderate to severe villus blunting, mild villus fusion, and moderate lymphoid depletion in the ileum. Table 1 summarizes the clinical and histopathological effects of the *S. Typhimurium* challenge in this porcine model. For representative photos of histopathological findings, see Supplemental Figure 1.

3.2 *S. Typhimurium challenge reduced ileal acetylcholine levels.*

The immune-modulatory role of the cholinergic system is mediated primarily through the ligand acetylcholine, so we asked whether bacterial enteritis would affect ileal mucosal acetylcholine concentration. Compared with controls, *S. Typhimurium* challenged animals exhibited a 3-fold reduction ($p=0.0142$) in ileal mucosal acetylcholine concentration (Fig. 1A).

3.3 *S. Typhimurium challenge down-regulates ileal mucosal acetylcholinesterase activity*

Since challenged animals demonstrated reduced concentrations of acetylcholine in ileal mucosa, we reasoned that AChE, the enzyme that degrades acetylcholine, might be

upregulated during enteritis. In contrast to what we expected, measurements for AChE enzymatic activity in ileal mucosa demonstrated over a 2-fold reduction in challenged animals compared with controls (Fig. 1B).

3.4 *S. Typhimurium* induced enhanced ChAT protein expression

Since alterations in AChE did not explain the reduction in acetylcholine concentrations, we next asked if the enzyme that produces acetylcholine, ChAT, was down-regulated in *S. Typhimurium*-challenged pigs compared to controls. Since there are several ChAT isoforms, we utilized Western blot to determine if there was a particular isoform that was increased following infectious enteritis. Again in contrast to our expected results, in ileal mucosa, we identified a ~60 KDa band that was increased in *S. Typhimurium*-challenged animals compared with controls (Fig 1C). This band corresponds to a ChAT isoform known as peripheral ChAT, which has been previously reported in porcine, non-human primate, rat, and guinea pig peripheral neurons, including the enteric nervous system. (Brehmer et al. 2004; Chiocchetti et al. 2003; Koga et al. 2013; Nakajima et al. 2000; Tooyama and Kimura 2000) No difference was found in the conical 80kDa ChAT isoform, commonly associated with the central nervous system, between challenged and control pigs (*data not shown*).

We next screened the mesenteric lymph nodes to determine if the change in cholinergic regulation extended beyond the gastrointestinal mucosa. In the mesenteric lymph nodes, a band similar in size and intensity to the ileum was found in *S. Typhimurium* challenged animals compared to controls (Fig 1D). In both the ileum and mesenteric lymph nodes, the expression of ChAT was significantly increased in challenged animals compared with controls (Fig 1C-D, respectively).

3.5 Cellular source of ChAT protein upregulation in *S. Typhimurium* challenged ileum

To gain further insight into the mechanism contributing to upregulated ChAT expression, we investigated the localization patterns of ChAT expression in ileal mucosa of control and challenged animals via immunohistochemistry. In agreement with Western blot analysis, ileum from *S. Typhimurium*-challenged pigs exhibited a larger area of ChAT expression (% Mucosa ChAT, Fig A-D, $p=0.06$). ChAT positive cells included villus and crypt epithelia (Figure 2E, F), lamina propria mononuclear cells (F) and submucosal and myenteric ganglia (not shown). Further, the % area of ChAT positive staining and intensity of staining was most pronounced within the epithelium of the Peyer's patch follicle associated mucosa (FAM) compared to follicle free mucosa (FFM) (Figure 2G-J).

3.6 Changes in cholinergic receptor gene expression following *S. Typhimurium* challenge.

Considering that *S. Typhimurium*-challenged pigs had reduced acetylcholine concentrations in ileal mucosa, we sought to determine if cholinergic receptor gene expression was altered. Cholinergic muscarinic receptors 1, 2, and 3 (CHRM1, CHRM2, and CHRM3, respectively) were selected due to their known prevalence in mediating inflammation (Hirota and McKay 2006a; Tobin et al. 2009). Utilizing real time - reverse transcription PCR with quantitation relative to control values, we determined that mRNA transcripts for CHRM1 and CHRM3 were significantly upregulated in ileal mucosa of pigs challenged with *S. Typhimurium* (Fig.

3). Specifically, CHRM1 expression was increased by ~ 4 fold compared with controls and CHRM3 expression was increased by ~2 fold relative to controls. There was no effect of *S. Typhimurium* challenge on CHRM2 expression (Fig. 3). We next measured mRNA expression of the cholinergic nicotinic alpha 7 receptor subunit (CHRNA7), which contributes to formation of the homopentameric receptor known to influence inflammatory responses (Matteoli et al. 2014; Wang et al. 2003). We found no effect of *S. Typhimurium* challenge on CHRNA7 expression in pig ileal mucosa (Fig. 3).

3.7 Correlation of mucosal ileal ChAT expression with clinical symptoms and histopathology

Since *S. Typhimurium* challenge influences several enzymes and receptors of the cholinergic system, we asked whether ileal mucosal ChAT expression was correlated with symptom severity and histopathological findings. Correlation analysis of ChAT expression intensity between rectal temperatures were performed separately for both controls and *S. Typhimurium* challenged animals. In controls, there was no correlation between ChAT expression and rectal temperature ($r=0.1363$, $p=0.8270$), (Fig 4A). However, in *S. Typhimurium* challenged pigs, a significant, positive correlation was found between ileum ChAT expression and rectal temperatures ($r=0.8985$, $p=0.0149$), (Fig. 4B).

Correlations between ileal mucosal ChAT expression and ileum histopathological scores for villus blunting, villus fusion, and lymphoid depletion were also performed separately for both controls and *S. Typhimurium* challenged animals. There was no villus blunting or villus fusion found in control pigs; therefore, no correlation could be performed (Fig 4C and 4E). While one control pig was found to have mild lymphoid depletion, overall there was no correlation between ileum ChAT expression and lymphoid depletion in controls was found ($r=0.1289$, $p=0.5528$), (Fig 4G). However, in *S. Typhimurium* challenged pigs, mucosal ChAT expression positively correlated with villus blunting ($r=0.9374$, $p=0.0185$) and villus fusion ($r=0.9366$, $p=0.0190$), (Fig 4D and 4F, respectively). Also, ileal mucosal ChAT expression had a trending positive correlation with lymphoid depletion; however, the relationship was not significant ($r=0.6317$, $p=0.1081$), (Fig 4H). A negative correlation was found between AChE and lymphoid depletion in *S. Typhimurium*-challenged animals ($r=-0.9296$, $p<0.0222$, data not shown).

3.8 Correlation of mucosal ileum ChAT expression with mucosal ileum cytokine expression

Given the positive correlation between ileal mucosal ChAT expression and rectal temperature and histopathology in *S. Typhimurium* challenged animals, we next asked if mucosal ChAT expression correlated with inflammatory cytokine production. No correlation between mucosal ChAT expression and TNF, IL-8, or myeloperoxidase (MPO) expression was found in controls (Fig 5A, 5C, 5E). However, in *S. Typhimurium* challenged pigs, a trend towards a positive correlation was found between ileum mucosal ChAT expression and ileum mucosal TNF expression, ($r=0.7354$, $p=0.0955$), (Fig 5B). No significant correlation was found between ChAT and IL-8 or MPO in *S. Typhimurium* challenged animals (Fig 5D and 5F). We also performed correlation analysis between acetylcholine concentrations or

AChE activity with ileum mucosal cytokine levels which showed no significant correlations (*data not shown*).

3. Discussion

4.1 *S. Typhimurium* challenge in pigs alters the expression of cholinergic proteins involved in ACh synthesis, receptor signaling and degradation

Results from the present study showed that acute oral *S. Typhimurium* challenge in pigs resulted in significant alterations in the expression of cholinergic system markers characterized by reduction in ileal mucosal ACh concentrations, down-regulation of acetylcholine degrading enzyme activity, AChE, and upregulation of acetylcholine synthesis enzyme, ChAT. Cholinergic muscarinic receptor 1 and 3 expression was also upregulated in *S. Typhimurium* challenged pigs.

To date, nematode-induced enteritis in rodents has been the most commonly utilized animal model to study pathogen-mediated modulation of the enteric cholinergic system, and many of these studies focused on the cholinergic myenteric plexus and motility. Commonly demonstrated across these studies, and very similar to our data, are that parasitic enteritis acutely induces reduced production and release of acetylcholine (Collins et al. 1989; Davis et al. 1998), elevated ChAT activity and expression (Davis et al. 1998; Palmer and Koch 1995), and diminished AChE activity (Davis et al. 1998; Palmer and Koch 1995) in the small intestine. Only a few studies exist reporting the impact of bacterial and viral infection on the enteric cholinergic system following chronic infections. In one study, *Helicobacter pylori* and Herpes Simplex Virus-1 infection were shown to induce chronic down-regulation of acetylcholine release, suggesting that like acute infection as observed in the present study, chronic infections may also act to suppress acetylcholine availability (Bercik et al. 2002; Brun et al. 2010). Similar to the acute infectious enteritis models, an acute chemical ileitis model of 2,4,6- trinitrobenzenesulfonic acid (TNBS) resulted in impaired ACh release with paradoxical increased ChAT expression when compared to controls (Vieira et al. 2017). Chemical colitis models utilizing TNBS and dextran sodium sulfate (DSS) have been shown to impact ChAT expression, though the effects were dichotomous, with TNBS reducing ChAT expression and DSS enhancing ChAT expression (Winston et al. 2013). In the present study, our results demonstrated similar paradoxical expression of impaired ACh production, with increased ChAT expression and reduced AChE function following acute bacterial enteritis. Therefore, previous reports and our data here suggest a highly conserved cholinergic response to acute intestinal injury across animal species and etiological agents.

4.2 Potential benefits and consequences of dynamic, enteric cholinergic expression changes during pathogen challenge

As mentioned above, the present study and previous work in rodent models support evidence for a conserved cholinergic response to intestinal injury characterized by a reduction in ACh levels and alterations in cholinergic system components involved in ACh synthesis and muscarinic receptor expression, and decreased ACh degradation. Together, this response may represent a compensatory mechanism by the host to preserve and recover cholinergic signaling in the face of diminished ACh levels. The functional significance of this response

is unclear at this time but might represent a beneficial response to limit intestinal mucosal injury. In support of this, cholinergic signaling through muscarinic receptor subtype 3 was shown to limit intestinal tissue damage in response to acute chemical colitis (Hirota and McKay 2006b). In the presence of inflammatory cytokines, TNF and INF- γ , muscarinic signaling preserved barrier function and limited release of neutrophil chemotactic cytokine IL-8, (Khan et al. 2015) demonstrating a protective role of cholinergic-mediated muscarinic signaling on the intestinal barrier potentially via control of the inflammatory milieu. Additionally, endotoxin induced upregulation of B lymphocyte ChAT expression was shown to negatively regulate neutrophil influx through action of acetylcholine on endothelium muscarinic receptors (Reardon et al. 2013). Likewise, down-regulation of mucosal AChE expression in response to endotoxin challenge was also shown to be necessary to limit acute cytokine production (Shaked et al. 2009). Increased cholinergic, muscarinic signaling, via muscarinic receptors may also play a role in adaptive immune responses. When challenged with enteric nematode infection, muscarinic receptor 3 knock out mice demonstrated an attenuated ability to generate humoral cytokines necessary to upregulate mucus production or alter motility, and this attenuated response resulted in increased worm and egg burden compared to wild type infected controls.(McLean et al. 2016) Further, muscarinic receptor 3 deficient mice infected with the enteric pathogen *Citrobacter rodentium* also demonstrated an impaired pathogen clearance (McLean et al. 2015) potentially linked with impaired adaptive immunity. Featuring a more direct role of muscarinic receptor signaling in adaptive immunity, muscarinic receptor 3 deficient lymphocytes demonstrated reduced cytokine production to cholinergic agonists with impaired memory immune responses and reduced activation and cytokine production in response to both humoral and cell mediated immunity inducing pathogens (Darby et al. 2015).

ACh can also bind and signal via α 7 nicotinic receptors, which have been more intensively studied in rodent chemical colitis models as major anti-inflammatory mechanism. In murine pathogen challenge models, α 7 nicotinic receptor genetic deficiency resulted in enhanced neutrophil recruitment which coincided with improved bacterial clearance compared with control animals (Giebelen et al. 2008). Pharmacological blockade of nicotinic receptors was shown to reduce intestinal villus damage induced by invasive *Shigella* infection (Svensson et al. 2004). In the present study, unlike muscarinic receptors, we did not observe significant changes in the expression of ileal α 7 nicotinic receptors in response to *S. Typhimurium* challenge. However, lack of changes in expression of α 7 nicotinic receptors following *S. Typhimurium* challenge does not necessarily imply an insignificant role for this receptor. For example, based on the preponderance of literature showing an inhibitory role of α 7 nicotinic receptors in immune activation, an unchanged expression level during pathogen challenge may facilitate bacterial clearance. Related to this, the reduced concentration of ACh in *S. Typhimurium* challenged pigs in the present may be an effort by the host to selectively activate muscarinic receptors over nicotinic receptors, since muscarinic receptors are known to have a much higher affinity for ACh compared with nicotinic receptors (Ahmad et al. 1987; Kellar et al. 1985; Purohit and Grosman 2006).

Together, results from our study with *S. Typhimurium* challenge in pigs and the literature support a critical role of the cholinergic response in balancing a robust immune responses required for pathogen clearance and development of adaptive immunity while at the same

time limiting excessive inflammation and loss of intestinal barrier integrity. Therefore, conditions that result in suppression or hyper-active enteric cholinergic tone could have significant consequences for host immunity and survival.

4.3 Increased ChAT expression and localization to the Peyer's patch epithelium in *S. Typhimurium* challenged pigs

Choline acetyltransferase (ChAT) has been shown to be expressed in neuronal and non-neuronal cell types. To gain more insight into the potential role of upregulated ileal ChAT expression in *S. Typhimurium*-challenged, investigated the localization of ChAT expression via IHC. As expected, we identified ChAT-positive cells within submucosal and myenteric neuronal ganglia and lamina propria which was increased in challenged pigs. Unexpectedly, we found the greatest intensity of ChAT-positive cells within the epithelium covering the Peyer's patch lymphoid mucosa. The mechanism for upregulation of ChAT, particularly in the Peyer's patch epithelium in challenged pigs remains to be determined. *S. Typhimurium* is an intracellular bacterial pathogen which establishes infection by transportation through microfold cells (M cells) over the Peyer's patches in the ileum or through invasion of absorptive ileum enterocytes (Broz et al. 2012; Meyerholz et al. 2002). After crossing the intestinal epithelial barrier, *S. Typhimurium* enters the lamina propria where the pathogen is phagocytosed by macrophage and dendritic cells of the immune system. The pathogen resides and replicates within these leukocytes, permitting dissemination to other organs such as the mesenteric lymph nodes (Broz et al. 2012; Rieger et al. 2015). Therefore, the focal pattern of ChAT localization observed in challenged pigs in present study follows the pathway of *S. Typhimurium* invasion. Others have shown that bacterial products such as LPS upregulate ChAT expression in mucosal associated lymphoid tissue, peritoneal B cells, macrophage, and dendritic cells (Kawashima et al. 2007; Reardon et al. 2013). Specifically in line with the current study, *S. Typhimurium* infection was shown to elicit induction of intestinal CD4⁺ Th17 lymphocytes, (Raffatellu et al. 2008) a group of cells that has recently been shown to be ChAT positive (Dhawan et al. 2016). Therefore we hypothesize that the specific ChAT localization patterns of Peyer's patch epithelium revealed in the present study is critical in the modulation of bacterial invasion and subsequent modulation of innate and adaptive immune response.

4.4 Summary

In summary, while the vagal cholinergic input has been shown to be an important key modulator of gut inflammatory responses, results from the present study demonstrate that the intrinsic cholinergic system, which includes neuronal and non-neuronal cell types, is dynamically regulated in response to *S. Typhimurium* challenge. The precise cause, cellular source, and significance of these cholinergic responses to overall host defense and immunity remains to be elucidated; however, our data establishes a foundation for future mechanistic research aiming to better understand the role of the enteric cholinergic system in mediating a balance between immune activation required for pathogen clearance and host immunity and the prevention of excessive inflammation and epithelial barrier damage.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Funding Sources

This work was supported by U.S. National Institutes of Health grants R01 HD072968 (to AJM), K08 DK097462 (to AJM), R03 DK097462 (to AJM). This project was also supported by Agriculture and Food Research Initiative Competitive Grant no. 2017–67015-26673 from the USDA National Institute of Food and Agriculture (to AJM).

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Highlights

- Typhimurium challenge in pigs resulted in a marked suppression of acetylcholine (ACh) levels in the ileum mucosa which coincided with a paradoxical upregulation of choline acetyltransferase (ChAT) and muscarinic receptor subtypes 1 (M1) and 3 (M3).
- Upregulated ChAT expression in *S. Typhimurium*-challenged pigs was localized predominantly in the ileal Peyer's patch epithelium.
- Ileal ChAT expression was positively correlated with rectal temperature, and histopathological severity in *S. Typhimurium* challenged pigs

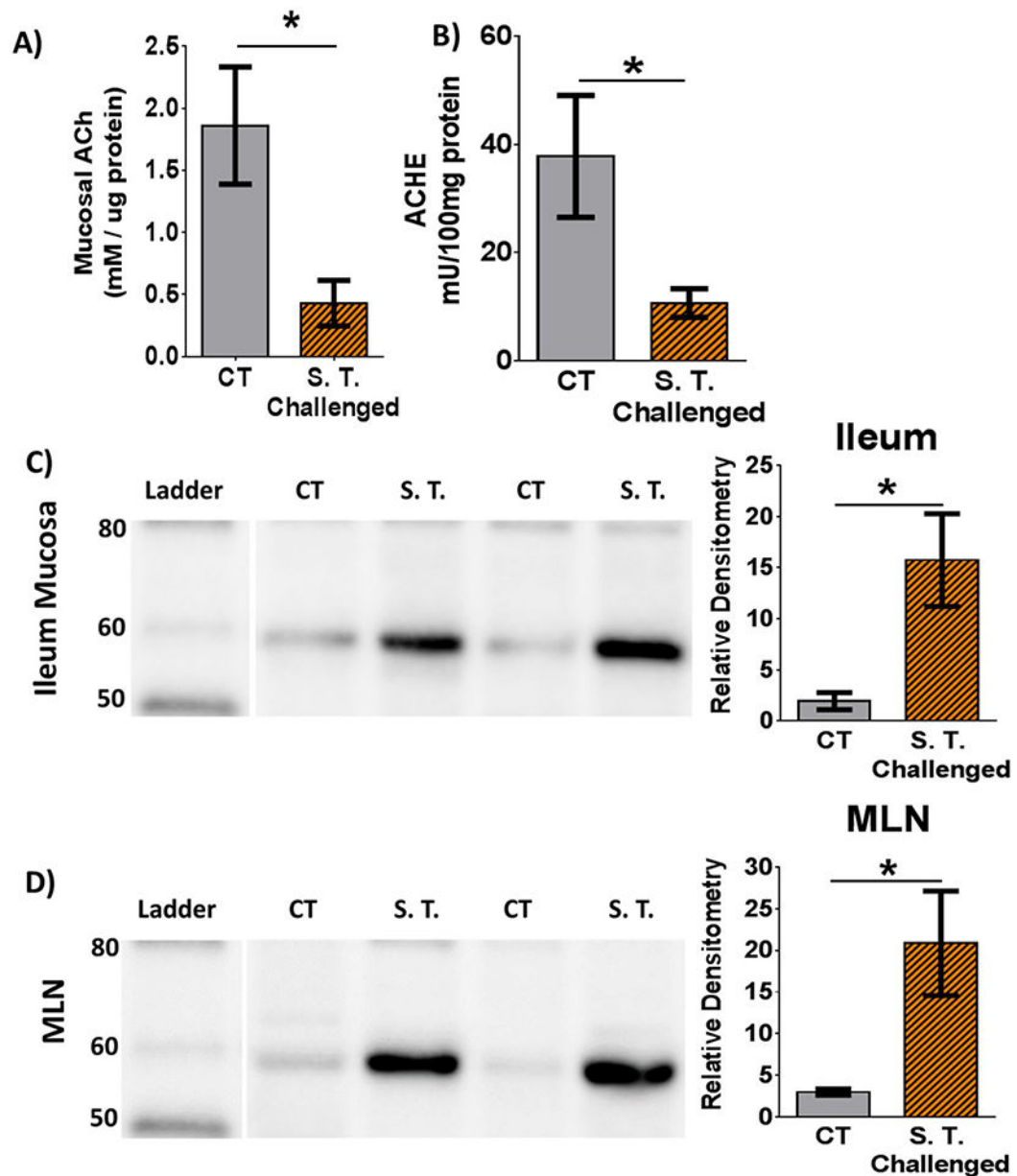


Figure 1. Impact of *S. Typhimurium* challenge on acetylcholine and cholinergic enzymes in ileum mucosa

A) Acetylcholine concentration in ileal mucosa. B) Acetylcholinesterase activity ileal mucosa. C) Western blot of ileal mucosa for porcine ChAT. Prominent band is between 60 and 50kDa. Histogram quantifying relative density directly to the right. D) Western blot of mesenteric lymph node (MLN) for porcine ChAT. Respective histograms represent relative density normalized to lane total protein. CT = Control, S.T. = *S. Typhimurium*. Bars and SEM represented. $n = 5$ controls and 6 challenged per bar. Student's t-test compared controls vs *S. Typhimurium* challenge. * = $p < 0.05$.

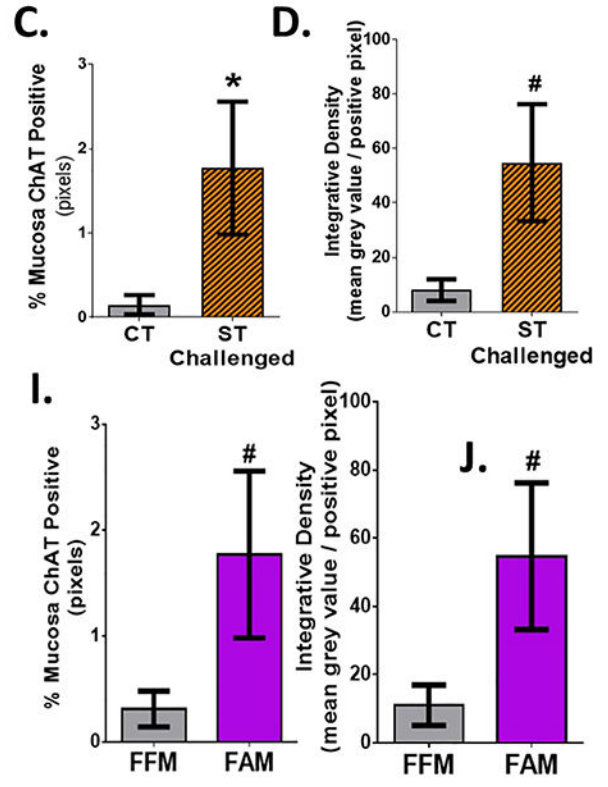
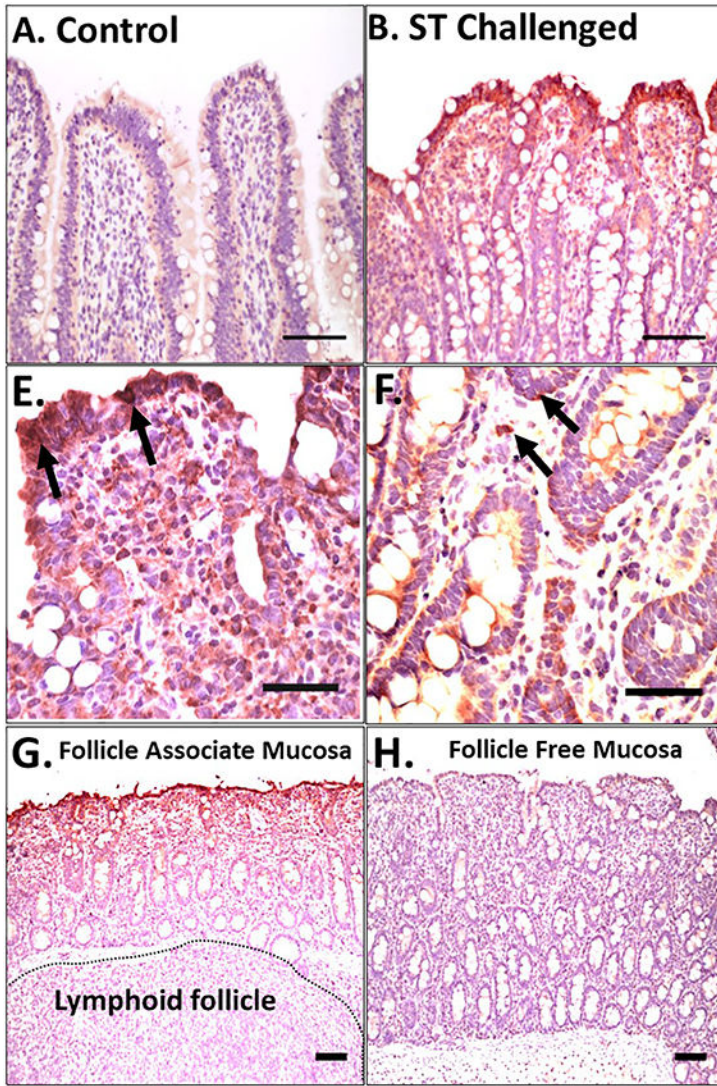


Figure 2. ChAT is elevated in epithelium and round cells of lamina propria over the Peyer’s patch following *S. Typhimurium* challenge
 A-B: 20x images of control (A) and ST challenged (B), scale bar = 100uM. C-D: % of area of ChAT positive mucosa (C) and integrated pixel density analysis (D) between control and ST challenged pigs. E,F: 40x representative images from ileum from an *S. Typhimurium* challenged animal demonstrating the enhanced epithelial expression (arrows) and staining within lamina propria monocytes (E,F; arrows). G-J: 10x image of (G) follicle free mucosa (FFM) and (H) follicle associated mucosa (FAM) . In (G), note the Peyer’s patch lymphoid follicle tissue directly under the mucosa. I,J: % area ChAT positive and integrated density between (I) FFM and (J) FAM .Scale bar - 100um. CT = Control, ST = *S. Typhimurium*. Mann-Whitney t-test was used in (C) and (G). Students T-test used in (D) and (H). # $p < 0.1$, * $p < 0.05$.

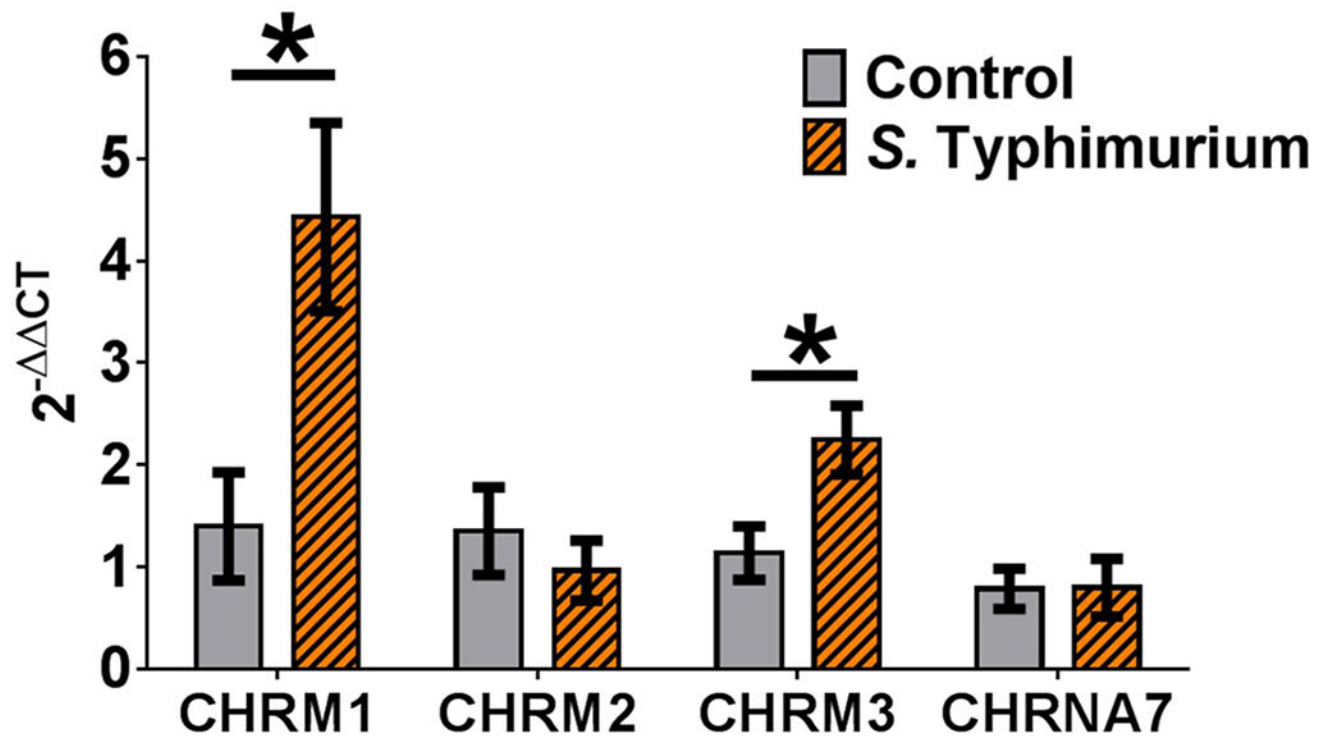


Figure 3. Impact of *S. Typhimurium* challenge on ileum mucosa cholinergic receptor gene expression

Salmonella Typhimurium enhances muscarinic receptor mRNA gene expression. mRNA gene expression was quantified utilizing two step reverse transcriptase quantitative PCR (RT-qPCR) for cholinergic muscarinic receptors 1, 2, 3 and cholinergic nicotinic receptor alpha 7 subunit (CHRM1, CHRM2, CHRM3, and CHRNA7 respectively). The data are presented as $2^{-\Delta\Delta CT}$ in gene expression relative to control group, normalized to *beta-actin* (*ACTB*) before statistical analysis. $n = 6$ controls and 6 challenged per bar. Student's t-test compared controls vs *S. Typhimurium* challenge of each cholinergic receptor. * = $p < 0.05$.

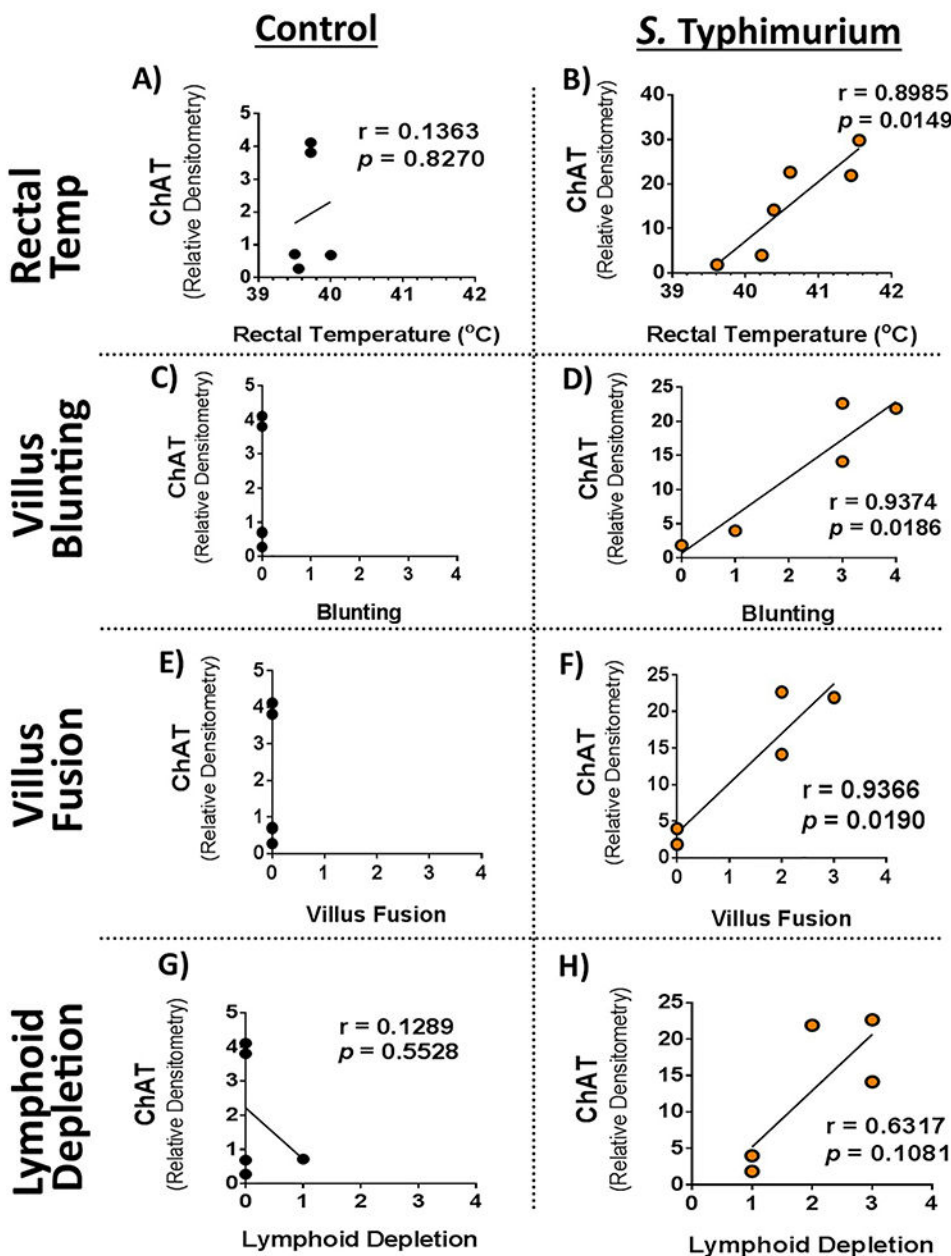


Figure 4. Correlation of ChAT with body temperature and ileum histopathology
 Rectal temperatures and histopathological scores were correlated with ileum mucosal ChAT expression from Figure 1. Correlations were performed independently on controls (A, C, E, G) and *S. Typhimurium* challenge (B, D, F, H) pigs. Pearson correlations were performed between mucosal ChAT and rectal temperature (A, B); Villus Blunting (C, D); Villus Fusion (E, F); and lymphoid depletion (G, H). Pearson correlation, r values and p values reported on figures.

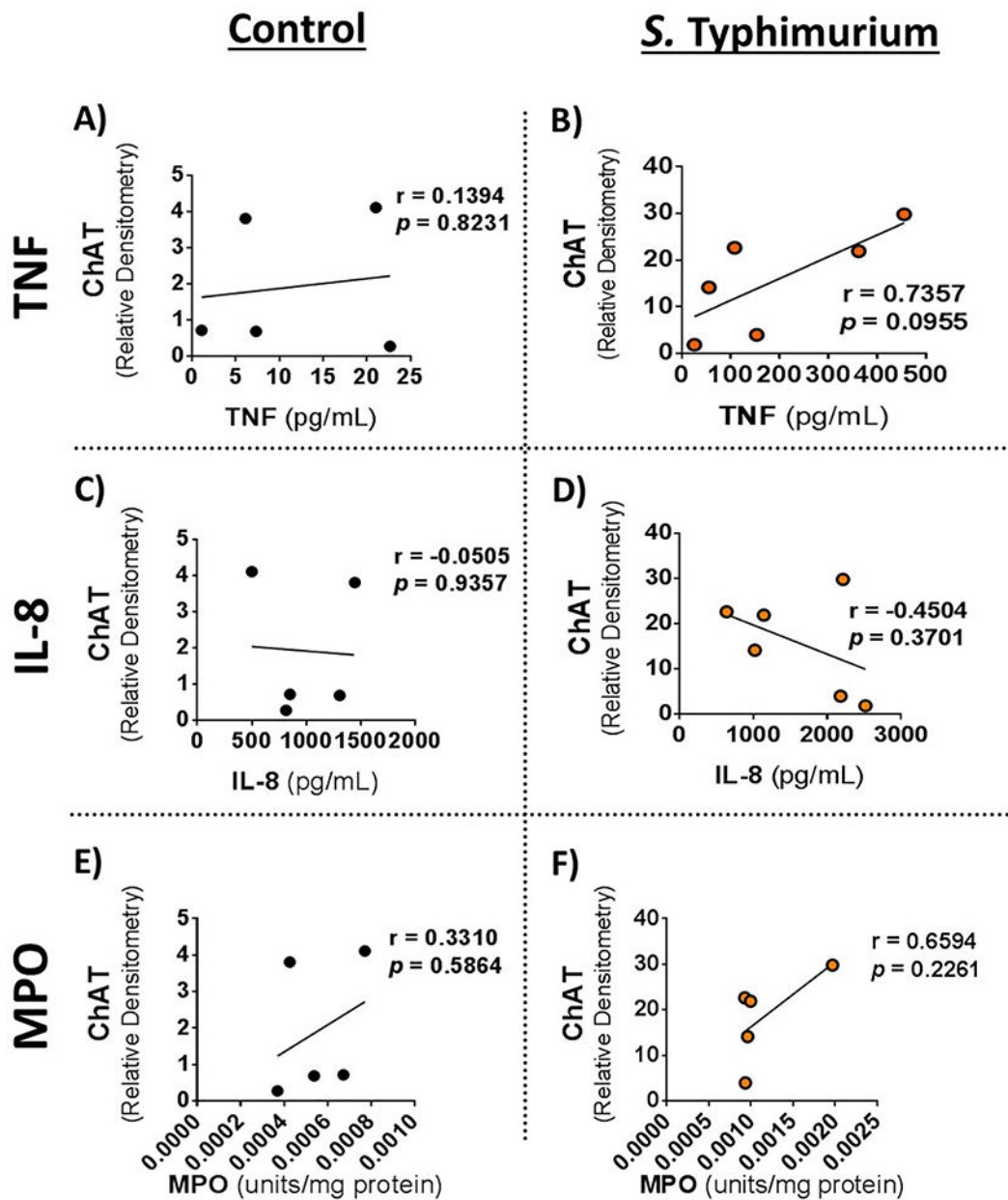


Figure 5. Correlation of ChAT with ileal mucosal cytokines
 Mucosal cytokine protein expressions were correlated with ileal mucosal ChAT expression. Ileal mucosal ChAT and cytokine correlations from A, C, E) controls and B, D, F) *Salmonella* Typhimurium challenge. ChAT correlation with mucosal TNF A, B); IL8 C,D); and MPO E,F). Cytokine protein levels were determined previously by ELISA and protein expression of ChAT, determined in Figure 1, from Western blot. Pearson correlation, r values and p values reported on figures. Each point represents an individual animal. R and p values for two tailed Pearson correlation were analyzed and are reported on each figure.

Table 1.
Previously Reported Clinical and Histopathological effects of *S. Typhimurium*

challenged Data in this table are summarized from a previously reported study.(Boyer et al. 2015) Fecal consistency was scored on a 4 point scale by an individual blinded to the study design (1= no diarrhea; 4= severe, profuse diarrhea). Histopathology scoring was performed as previously reported. Villus blunting was scored on a 5 point scale with 0 = 1:4 crypt to villus height and 4 = no villus present. Villus fusion and lymphoid depletion were scored on a 4 point scale with 0 = normal and 3 = severe. n=5–6 per group. *p*values generated by two tailed students t-test. For significance, y=yes; n=no.

	Control	<i>S. typhimurium</i> challenged	P value	
Fecal Score	1.20	3.33	0.0012	
Rectal Temperature(°C)	39.70	40.64	0.0238	
Histopathology Scores	Villus Blunting	0.00	2.20	0.0172
	Villus Fusion	0.00	1.40	0.0479
	Lymphoid Depletion	0.20	2.00	0.0063