***Odoribacter splanchnicus* mitigates *Salmonella*-induced gut inflammation and its**
associated pathogenesis via its secreted bacteriocin

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ABSTRACT

Foodborne pathogens continue to be a leading concern of health hazards worldwide claiming

the lives of millions. The emergence of drug-resistant strains poses a great threat to global

world health management. Therefore, designing novel therapeutic strategies aiming to

overcome pathogen burden is of utmost importance. Here, we identified the crucial role of a

gut-commensal species, *Odoribacter splanchnicus* in mitigating *Salmonella* pathogenesis in

mice by inhibiting gut vascular barrier disruption, acute inflammatory infection signs, *in vivo*

biofilm formation and by preserving tight junction protein functions upon pre-colonization.

Further, our *in vitro* studies revealed that not only live *O. splanchnicus* (OS) but also its culture

supernatant inhibit *Salmonella* biofilm formation, intracellular proliferation in human intestinal

cells, and its virulence gene expression. In addition, OS's inhibitory effect on *Salmonella* is specific as *Enterococcus faecalis* fails to exert an inhibitory effect on *Salmonella*. Further, our results depicted that the specific protective role of OS acts over a broad spectrum as it confers protection against flagellated Gram-positive, *Listeria monocytogenes* and Gram-negative, *Salmonella* Typhimurium foodborne pathogens, as opposed to non-flagellated *Shigella flexneri*. Moreover, OS conferred protection even after its administration to mice post-establishment of infection highlighting its therapeutic potential. Using several biochemical and proteomics approaches, we characterized the key active molecule secreted by OS to limit intracellular *Salmonella* and *Listeria* replication in human intestinal epithelial cells by regulating key virulence effectors and flagella. Collectively, our study highlights the broad-spectrum protective role of OS in mitigating *Salmonella* and *Listeria* pathogenesis and implicates its therapeutic potential.

Keywords: *Salmonella* pathogenesis, **Gut vascular barrier permeability**, **inflammation**, **inter-species bacterial regulation**, ***Listeria***, ***Odoribacter splanchnicus***, **gut commensal**, **biofilm, flagella**

INTRODUCTION

The existence of the gut vascular barrier (GVB) acts as a protective barrier against the entry of microorganisms into systemic blood circulation[1, 2]. Enteric pathogens such as *Salmonella enterica* can cross the mucus and intestinal epithelial barrier, ultimately leading to GVB damage and systemic dissemination[1]. The gut microbiota plays another major role in the maintenance of the intestinal epithelial barrier by the production of several metabolites such as Short-chain fatty acids (SCFAs), indoles, and polyphenol metabolites that contribute to epithelial barrier integrity and turnover by regulating the expression of Tight-junction (TJ) genes[3]. Additionally, these SCFAs have the potential to exert anti-inflammatory properties by regulating cytokine production, mucosal immune cell migration, and secretion of

antimicrobial peptides (AMPs) by intestinal epithelial cells and macrophages(4, 5). Our

previous study reported an incidence of increased *Salmonella* dissemination in blood, liver, and

spleen upon SIRT1 and SIRT3 inhibition owing to the increased inflammatory response(6).

Salmonella gains entry inside the body through the orofecal route along with the ingestion of

contaminated food and water. *Salmonella* resist the low acidic pH of the stomach by utilizing

the acid tolerance response (ATR) genes and finally reach the lumen of the small intestine(7).

In the intestinal tract, it uses specialised microfold (M) cells for macro-pinocytic entry. The

pathogen gains entry into the non-phagocytic epithelial cells via its *Salmonella* Pathogenicity

Island-1 (SPI-1) effectors to breach the intestinal barrier (8). Once traversing through the

mucosal lining of the gut epithelium, it reaches the lamina propria (8). Apart from M cells, the

dendritic cells can also sample the bacteria from the lumen to the lamina propria. In the lamina

propria, it encounters various host immune cells like neutrophils, macrophages, dendritic cells

and so on. Macrophages, dendritic cells, and neutrophils are responsible for successful

dissemination throughout the body through the reticulo-endothelial system (RES) to the

secondary organs such as Mesenteric Lymph Node (MLN), liver and spleen(8). *Salmonella*

relies on the SPI-2 secretion for its intracellular and intravacuolar life by injecting various

virulence factors into the host cell cytosol (9).

Here, in this present study, we aimed to understand this exact mechanism of *Salmonella*

dissemination and whether colonization with a gut commensal like *Odoribacter splanchnicus*

(OS) could ameliorate the *Salmonella*-induced inflammation and in turn its pathogenic

dissemination. OS, belonging to the order Bacteroidales, is a common, member of the human

intestinal microbiota. A decrease in *Odoribacter* abundance has been linked to different

microbiota-associated diseases, such as non-alcoholic fatty liver disease, cystic fibrosis and

inflammatory bowel disease (IBD)(10). A recent report indicated the ability of OS and its

effectors transported in Outer membrane vesicles (OMVs) to exhibit anti-inflammatory effects

on the gut epithelium(10). Further, OS has been implicated to confer resistance against colitis

and colorectal cancer(11).

Our study shows that OS colonization reduces *Salmonella* gut colonization, its organ load, and

its systemic dissemination even under aggravated inflammatory responses such as SIRT1 or

SIRT3-inhibited condition. OS ameliorates acute *Salmonella*-induced inflammation and its

associated pathogenesis by inhibiting GVB damage and preserving tight junction protein

functions. Further, we demonstrate that not only live OS but also its culture supernatant inhibits

Salmonella in vitro biofilm formation, intracellular proliferation in human intestinal cells, and

its virulence gene expression. Subsequently, our results depict that the specific protective role

of OS acts over a broad spectrum as it confers protection against both flagellated Gram-

positive, *Listeria monocytogenes* and Gram-negative, *Salmonella* Typhimurium foodborne

pathogens in contrast to non-flagellated *Shigella flexneri*. Furthermore, OS conferred

protection even after its administration to mice post-establishment of infection highlighting its

therapeutic potential. Subsequently, we identified the key OS-secreted bacteriocin protein

responsible for limiting intracellular *Salmonella* and *Listeria* replication in human intestinal

epithelial cells by regulating virulence genes such as flagella. Overall, our study implicates the

broad-spectrum protective role of OS in mitigating foodborne infections caused by *Salmonella*

and *Listeria* pathogenesis.

RESULTS

***O. splanchnicus* hinders *Salmonella* colonization in mouse gut and its systemic**

dissemination

To denote the role of OS in governing *Salmonella* pathogenesis, we colonized C57BL/6 mice

with 10^5 CFU of OS 1 day prior to *Salmonella* infection via the oral route and validated the

colonization by undertaking OS 16S rRNA qPCR analysis (Fig 1, A-B). OS pre-colonized mice

showed decreased susceptibility to *Salmonella* infection in comparison to the non-colonized mice cohort which succumbed to death by day 7 post-infection (p.i.). Contrastingly, 100% of the OS colonized mice were alive even at day 8 post *Salmonella* infection (Fig 1, C). OS non-colonized mice exhibited increased weight loss percentage upon *Salmonella* infection and OS pre-colonization led to diminished weight reduction percentage (Fig 1, D). The increased *Salmonella* organ loads and bacterial dissemination in the non-colonized mice groups were reversed upon OS colonization (Fig 1, E-F). Our previous findings revealed increased bacterial dissemination owing to heightened inflammatory responses in the case of SIRT1 and SIRT3 knockdown or inhibitor-treated conditions(6). Therefore, in this study, we intended to examine whether OS could exert its protective role against *Salmonella* infection even in the case of aggravated inflammatory conditions such as in the case of SIRT1 or SIRT3 inhibited conditions. Our result suggests that even in aggravated conditions of inflammation and *Salmonella* dissemination under SIRT1 or SIRT3 inhibitory conditions, OS could prevent *Salmonella* colonization and its dissemination (Fig 1,D-G).

O. splanchnicus colonization ameliorate the acute *Salmonella-induced* inflammation and its associated pathogenesis

Heightened inflammatory responses facilitate *Salmonella* dissemination in mice(6). Thus, we hypothesized that OS gut colonization might ameliorate the *Salmonella*-induced inflammation. ELISA-mediated IL-6 serum quantification revealed reduced IL-6 production upon *S.* Typhimurium (STM) infection in the OS colonized mice group to the non-colonized cohort (Fig 2, A). The reduced IL-6 production upon OS colonization was even observed in elevated inflammatory conditions of SIRT1 or SIRT3 inhibition. To combat the enteric pathogen challenge, the host employs one of its major defence arsenals, that is, the production of antimicrobial peptides (AMPs)(12, 13). STM infection induces enhanced mRNA expression of AMPs in mice with respect to the uninfected control. We found that OS colonization in mice

resulted in the downregulation of several AMP genes such as Lysozyme-C1 precursor,

Angiogenin-4 precursor, CCL20, Beta-defensin-4 and Regenerating-islet-derived 3 beta-

precursor (Fig 2, B). Mucus lining serves as another crucial barrier against the entry of enteric

pathogens. However, enteric pathogens adopt several strategies to circumvent this barrier to

reach the underlying epithelium. To assess the mucin layer dynamics upon the *Salmonella*

challenge, we assessed MUC2 expression in the ileal tissues of OS colonized versus non-

colonized mice groups. Our result depicted enriched MUC2 expression in the OS non-

colonized mice cohort compared to the colonized cohort (Fig.2, C). The increase in the mucus

lining upon *Salmonella* infection could be one of the host defense mechanisms to bolster the

barrier and exclude pathogens(14). As per previous literature, the rate of mucin production is

modulated by several stimuli including inflammatory cytokines (15), vasoactive antimicrobial

peptides, and other innate and adaptive immune responses (16). Germ-free mice depict a

reduced number of goblet cells, and mucin storage granules(17) as well as reduced expression

of AMPs(18). The haematoxylin eosin-stained intestinal sections suggested increased intestinal

tissue damage with irregularly organized villi, necrotic submucosa layer, and detached

intestinal cells in all the STM-infected non-OS-colonized cohorts. However, OS pre-

colonization restored the intestinal tissue architecture in all the mice cohorts (Fig 2, D).

O. splanchnicus restricts _Salmonella_-mediated GVB damage by promoting tight junction

gene expression

Salmonella enters the intestinal epithelial milieu by breaching the gut vascular barrier (GVB)

and by inducing tight junction disruption. Upon ingestion, *Salmonella* like other food-borne

antigens enters into the Mesenteric Lymph Node (MLN) by accessing the lymphatics and

subsequently gains entry into the liver via the bloodstream(19). The translocation of the

pathogen across the blood endothelial cells is governed by the integrity of the GVB(1). The

GVB architecture is maintained by several tight junction and adherens junction proteins that

control paracellular trafficking(20, 21). PV-1 (plasmalemma-vesicle associated protein-1) expression is a determinant of GVB damage and heightened PV-1 expression denotes EC permeability(22, 23). Our data depict heightened PV-1 marker expression alongside reduced tight junction protein mRNA expression such as *Tjp1* and *Claudin5* upon *Salmonella* infection correlating with increased bacterial dissemination in blood and liver (Fig. 3, A-B, Fig. S1, A). CD34, an endothelial marker lining the blood vessels (24, 25), has been promoted upon OS colonization in *Salmonella*-infected mice intestines indicative of intact vasculature (Fig. 3, B-D, Fig.S1, B). On the contrary, OS mice cohorts exhibited reduced CD34 expression in secondary sites of dissemination such as the liver, suggesting toward reduced vasculature (Fig.S1, C). Therefore, OS pre-colonization attenuates the *Salmonella* dissemination by exhibiting reduced PV-1 protein expression and promoting tight junction protein expression (Fig. 3, A-D). Angiogenesis is one of the processes that facilitate bacterial dissemination. Previous studies in *Mycobacterium marinum* have shown that transcriptional induction of pro-angiogenic markers like *Vegfa* contributes to increased bacterial burden and dissemination(26). In our study, OS pre-colonization in the gut resulted in reduced expression of the pro-angiogenic *Vegfa* in the infected liver tissues, which coincided with the reduced *S.* Typhimurium burden in the bloodstream and liver (Fig S1, D). Together, our results depict the ability of OS to suppress angiogenesis and in turn STM dissemination. OS could exert its angiogenesis inhibitory effects even in instances of aggravated dissemination conditions like in the case of SIRT1 and SIRT3 inhibitor treatment in mice. Previous literature suggested reduced expression of *Axin2,* a β catenin target gene, upon wildtype *Salmonella* infection, even after Wnt3a stimulation suggesting that *Salmonella* affects the GVB by inhibiting canonical Wnt signalling (1). Here, our results suggest that OS pre-colonization restores the *Salmonella*-triggered attenuated Wnt signalling response with increased Axin2 expression in the *Salmonella*-infected intestinal and thereby restricts GVB damage and dissemination (Fig. S1,

E). The integrity of the infected mice gut was estimated by the gut permeability assay using

FITC-Dextran. Our results indicate increased gut vascular permeability upon *Salmonella*

challenge which declined upon OS pre-colonisation.

***O. splanchnicus* culture supernatant restricts *Salmonella* biofilm formation and**

intracellular proliferation in human intestinal cells

To delineate the mechanism behind the protective role of OS against *Salmonella* pathogenesis,

we performed *Salmonella* infection studies in human intestinal epithelial cell line Caco-2 in

the presence of OS. The intracellular survival assay suggested decreased STM invasion and

subsequently attenuated intracellular replication within the infected Caco-2 cells upon co-

infection with live OS and in the presence of OS culture supernatant (Fig 4, A-B). However,

STM co-infection with fixed OS showed percent invasion and intracellular replication like that

of STM infection (Fig 4.5, A-B). Together, these results hinted at the presence of some secreted

inhibitory components in the OS culture supernatant that has the propensity to limit STM

infection. The reduced inhibition and proliferation of STM within Caco-2 cells in presence of

OS was subsequently validated in immunofluorescence studies, wherein STM showed a

significant reduction in invasion at 2hr p.i. and intracellular proliferation at 16hr p.i. in case of

OS co-infection (Fig 4, C-D). Similar results were obtained upon infection with the human-

restricted serovar *S.* Typhi (STY) to the Caco-2 cells wherein OS and its supernatant attenuated

STY invasion and intracellular replication (Fig. 4, E-F). Further, we assessed the *in vitro* growth

of the STM in co-culture conditions with OS or in the presence of OS culture supernatant in

anaerobic conditions to mimic the microaerophilic or hypoxic environment of the gut or in

aerobic culture conditions. We observed no significant delay *in vitro* growth of STM at its

initial lag to log phase of its growth in the presence of OS supernatant or with live OS under

anaerobic culture conditions (Fig S2, A-B). In aerobic growth conditions as well, *Salmonella*

growth remained unperturbed even in the presence of OS or its culture supernatant (Fig S2, C-

D). Several previous studies have corroborated the ability of commensal bacterial culture supernatants to inhibit colonization and the biofilm formation of several pathogens. For instance, a commensal bacterium, *Staphylococcus epidermis*, could inhibit biofilm formation and nasal colonization of its pathogenic counterpart, *Staphylococcus aureus* (27). Moreover, biofilm formation is also regulated at the interspecies level. Indole is an inter-species biofilm regulator wherein it decreases *E. coli* biofilm formation and facilitates biofilm formation of pseudomonads (28). STM form biofilm to colonize host, evade host defence arsenals and antibiotics, persist in case of chronic infections, and transmit to new hosts (29, 30). We, therefore, examined the biofilm formation of STM in presence or absence of live, PFA-fixed-OS or OS culture supernatant. STM exhibited biofilm formation defect, in the presence of live OS or OS supernatant (Fig 4, G). However, the STM biofilm inhibitory property of STM was diminished upon treatment with paraformaldehyde (PFA)-fixed OS (Fig 4.5, G). Further, qPCR-mediated gene expression studies of *Salmonella* biofilm genes revealed decreased expression of genes regulating biofilm (*csgD*, *csgA*, *bapA*) under treatment with live OS or its culture supernatant (Fig.4, H-J). Flagella-mediated attachment and several virulence factors such as SPI-1 and SPI-2 genes are responsible for *Salmonella* pathogenesis and its biofilm formation (31-33). Therefore, we evaluated a few of the *Salmonella* flagellar and SPI-1 and SPI-2 gene expression in the presence or absence of OS or its culture supernatant via quantitative real-time PCR. Our results depicted attenuated expression of flagellar gene (*fliC*), SPI-1 (*sipC*) and SPI-2 (*pipB2*) genes both in the presence of live OS and its spent media but not in presence of PFA fixed-OS (Fig. 4, K-M). The loss in *Salmonella* flagellar motility was further validated via TEM imaging wherein OS or its supernatant reduced the flagella formation of *Salmonella* considerably (Fig.4, N). Together, our results implicate the ability of OS to inhibit STM biofilm formation via active secretion of certain inhibitory molecules in the supernatant.

O. splanchnicus inhibitory effect on Salmonella is specific as Enterococcus faecalis, another gut resident microbe, fail to inhibit Salmonella intracellular proliferation or biofilm formation

The resident gut microbes provide colonization resistance against invading pathogens by deploying several mechanisms, including the production of inhibitory molecules, nutrient competition, maintenance of the intestinal epithelial integrity, immune activation, and via the deployment of bacteriophages (34, 35). To understand whether our observed inhibitory trait of OS against *Salmonella* is due to a common colonization restriction phenomenon of gut microbes via competitive inhibition, or it is a specific phenomenon, we evaluated the role of another gut resident microbe, *Enterococcus faecalis* on *Salmonella* intracellular replication in human intestinal epithelial cells and *Salmonella* biofilm formation. *Enterococcus sp.* belong to the phylum of Firmicutes in the family Enterococcaceae. Approximately, 10^6-10^7 *Enterococcus sp*. reside in the human intestine with major abundance of *E. faecalis* (10^5-10^7 CFU/gm faeces) and *E. faecium* (10^4-10^5/gm faeces) (36). *E. faecalis*, *E. faecium*, *Bifidobacterium sp. Lactobacillus sp.,* and *E. coli* colonize the gut of healthy breastfed infants during the initial 7-10 days post birth (37). Our intracellular survival assay data showed no defect in *Salmonella* invasion or intracellular replication capability in Caco-2 cells in presence or absence of *E. faecalis*, its culture supernatant or PFA-fixed *E. faecalis* (Fig. 5, A-B). Moreover, *E. faecalis* or its *E. faecalis* supernatant did not inhibit *Salmonella* biofilm formation (Fig. 5, C). Together, our results suggest the inability of *E. faecalis* to inhibit *Salmonella* invasion, intracellular proliferation or its *in vitro* biofilm formation, thereby highlighting the specificity of OS in restricting *Salmonella* pathogenesis.

O. splanchnicus confers protection against another foodborne pathogen Listeria monocytogenes but not against Shigella flexneri

To understand whether OS exerts a common broad-spectrum protection or colonization

resistance against other foodborne pathogens such as *Listeria monocytogenes* or *Shigella*

flexneri, we orally infected 6-8-week-old C57Bl/6 mice with 10^8 or 10^7 CFU of *L.*

monocytogenes (LM) or *S. flexneri* (SF)*,* respectively 1 day post colonization with 10^5 CFU of

O. splanchnicus for organ burden evaluation. For survival study, mice were orally gavaged

with 10^9 or 10^8 CFU of *L. monocytogenes* or *S. flexneri,* respectively. OS-colonized mice

exhibited increased survival rates upon LM challenge in comparison to the non-colonized mice

cohort (Fig. 6,A). Contrastingly, both OS-colonized and non-colonized mice showed

comparable survival upon SF challenge (Fig. 6, G). In line with the survival data, increased

bacterial burden was observed in blood, MLN, spleen and liver alongside enhanced percent

loss in mice body weight, increased GVB damage, mucin and AMPs production in the case of

Listeria infection in non-colonized mice group with respect to OS-precolonized- mice group

(Fig. 6, B-F). Conversely, *Shigella* infection resulted in comparable organ loads and

comparable signs of pathogenesis in both cohorts (Fig. 6, H-L). In a similar line to *Salmonella*

infection, OS mitigated host antimicrobial response with reduced expression of antimicrobial

peptide transcripts and Muc2 production during LM infection as opposed to SF infection

amounting to reduced intestinal tissue damage. Further, OS promoted tight junction functions

during *Listeria* infection with reduced GVB damage and *Vegf* expression unlike during *Shigella*

challenge (Fig. S3, A-J). Moreover, OS colonization could restrict the *in vivo Salmonella* and

Listeria biofilm within the spleens of the infected mice cohorts as opposed to the non-colonized

mice cohorts (Fig.S3, K-L). The contrasting results between *Listeria* and *Shigella* could be

attributed to the presence of flagella in both *Salmonella* and *Listeria* but not in *Shigella*. Like

Salmonella, *in vitro* culture of *Listeria* with OS or its culture supernatant led to a loss in flagella

as evidenced via the TEM imaging (Fig.S3, M). Since flagella is one of the factors facilitating

biofilm formation and invasion, OS could inhibit both *Salmonella* and *Listeria* colonisation

and biofilm formation. Similar to *Salmonella* infection, OS colonization enhanced CD34 expression in *Listeria*-infected mice intestines and reduced CD34 expression in the *Listeria*-infected liver (Fig. S3 N-O). Further, LM exhibited attenuated intracellular proliferation within human intestinal Caco-2 cells upon live OS or OS supernatant treatment (Fig. S4). Altogether, this result hints at the potency of OS to act broadly against Gram-positive and Gram-negative bacteria and probably via regulation of flagella formation.

The protective role of *O. splanchnicus* against *Salmonella* Typhimurium and *Listeria* monocytogenes persists even on administration post-establishment of infection

Next, we sought to examine whether the protective role of OS against *Salmonella* and *Listeria* is sustained when the infection has been fully developed and whether it can serve as a treatment regimen against foodborne diseases. Our results indicate the protective ability of OS against both foodborne pathogens in their intestinal phase of infection when administered at 1-day post-infection (dpi) as depicted by lower pathogen loads in all the organs and blood along with reduced percent weight loss at 5th-day post-*Salmonella* or *Listeria* challenge in the OS-administered (1-dpi) cohort when compared to the untreated cohorts (Fig. 7,A-K). Although the protective ability of OS persists even upon establishment of infection when administered at 3rd-day post-*Salmonella* or *Listeria* challenge as indicated by reduced organ loads and percent weight loss in the OS-colonized cohorts, the extent of protection is lower compared to its administration at 1-dpi (Fig. 7, L-W), Further, the OS challenge at 1-dpi reduced the mice's susceptibility to the *Salmonella* and *Listeria* challenge(Fig. 7, L,R). However, administration of OS at 3-dpi delayed the death of the *Salmonella* and *Listeria*-infected mice cohorts by 1 day (Fig. 7, L,R). In line with the survival and organ burden data, OS treatment at both 1st and 3rd dpi reduced both *Salmonella* and *Listeria*-induced GVB damage with reduced PV-1 expression, mucin, AMPs, and *Vegf* expression, alongside promoting tight junction protein (*Tjp1* and *Claudin5*) and *Axin2* expression (Fig. S5A-N). However, OS administration at 3-dpi conferred

less protection than at 1-dpi as indicated by the protein expression profiles of Muc2, PV-1, and tight junction protein ZO-1 (Fig. S5, O-P). Moreover, the hematoxylin- eosin-stained intestinal sections revealed reduced intestinal tissue damage with restoration in the villi architecture in the 1- and 3-dpi OS-administered mice cohorts compared to the non-OS-treated groups (Fig.7, X).

Moreover, OS colonization could restrict the *in vivo Salmonella* and *Listeria* biofilm within the spleens of the infected mice cohorts as opposed to the non-colonized mice cohorts both when administered at 1-day and 3-day post-pathogen challenge. (Fig.7, Y-Z). Additionally, OS colonization mitigated gut vascular permeability during *Salmonella* and *Listeria* challenges as indicated by reduced FITC-dextran fluorescence units in the plasma both upon pre or post-OS colonization.

Furthermore, OS alleviated *Salmonella*-induced intestinal inflammation with reduced IL-1β production in the infected intestinal tissues both upon pre-colonization and post-colonization at 1-dpi (Fig. S6).

Characterization of the inhibitory active molecule within *O. splanchnicus* culture supernatant

Our results suggested that not only live OS but also its spent culture supernatant could inhibit *Salmonella* invasion and intracellular proliferation in intestinal human epithelial cells. Moreover, the OS and its supernatant could inhibit STM biofilm formation *in vitro*. This inhibitory effect disappeared upon co-culture with paraformaldehyde-fixed-OS indicating the presumable role of certain inhibitory microbial products secreted by metabolically live OS. To delineate the nature of the secreted molecule/s, we subjected the supernatant to various conditions such as high temperature, divalent cationic chelator, EDTA, and protease (proteinase K), and subsequently assessed the biofilm formation (Fig. S7, A-C). Our biofilm assay

suggested that the active molecule is resistant to EDTA treatment and is stable at higher

temperatures (Fig. S7, A-B). Nevertheless, the active molecules within the fraction were

sensitive to the proteinase K treatment, suggesting the possibility of being protein in nature

(Fig. S7, C). SDS-PAGE of the OS supernatant revealed the size of the differential proteins

with respect to the media control ranged below 65kDa. (Fig. S7, D). Concentration-dependent

estimation of the biofilm activity of the supernatant revealed that the active molecules retain

their activity at a minimal concentration of 0.4% v/v (Fig. S7, E). We further fractionated the

supernatant using an Amicon 3k MWCO and 10k MWCO ultra filter device (Fig. S7, F-G). We

found that the active molecule/s is/ are higher than 3kDa cut-off range eliminating the role of

small molecules and ions (Fig. S7, F-G). A summary of the conditions tested for active

molecule characterisation and their effective outcome has been summarized in Fig. S7, H.

Subsequently, we subjected the culture supernatant to gel filtration chromatography (Fig. S7,

I). Out of all the fractions tested, around 24 fractions showed a biofilm inhibitory role. These

24 fractions were further subjected to a titration experiment to zero down upon the three

fractions with the highest inhibitory activity (A82,A83,A84) (S7, J-L). The gel filtration

standard curve run suggested that the active molecules fall in the size range of <60kDa and

>3kDa (Fig. S7, I). These three fractions along with the total concentrated supernatant (serving

as positive control) and fractions with no or least inhibitory activity (serving as negative

control-A31,B12) were ultimately subjected to Reverse Phase LC-MS mass spectrometry.

Upon analysis of the proteomics dataset (Fig. S7, M), we obtained 364 proteins showing

enrichment in the positive fractions. Among the 364 proteins, 74 proteins exhibited known or

predicted function as per UniProt and STRING database and 11 proteins were linked to

antibacterial, bacteriostatic, flagella, and biofilm regulation based on data curation and

literature survey (Table-S1). We selected bacteriocin proteins exhibiting direct association with

antibacterial activity for further analysis. These proteins include bacteriocins which are

antimicrobial peptides that can kill or inhibit closely related or unrelated bacterial strains(38,

39).

O. splanchnicus*-derived purified peptide, bacteriocin, limits both *Salmonella* and *Listeria

pathogenesis

Treatment of the Caco-2 cells with the purified bacteriocin inhibited STM, STY, and LM

invasion and fold proliferation without compromising host-cell viability (Fig. 8, A-D).

Additionally, the purified bacteriocin treatment inhibited STM biofilm formation in a dose-

dependent manner alongside reduced transcript level expression of *Salmonella* biofilm

regulatory genes (Fig. S8, A-D). Moreover, bacteriocin treatment (10µg/ml) attenuated

Salmonella virulence gene expression such as SPI-1 gene- *sipC*, SPI-2 gene-*pipB2,* and

flagellar gene *fliC* (Fig. S8, E-G). Subsequently, TEM imaging revealed a complete loss of

Listeria flagella upon treatment with the OS-derived bacteriocin in contrast to *Salmonella*

which retained few flagella even on bacteriocin treatment (Fig. 8, E-H). Furthermore, the

motility assay demonstrated a significant swimming defect of *Listeria* and *Salmonella* upon

treatment with either OS supernatant or bacteriocin to the untreated control (Fig. 8, I-J).

Overall, our study depicts the novel role of OS-derived bacteriocin in limiting *Salmonella* and

Listeria invasion and pathogenesis by regulating virulence effectors such as SPI-1, SPI-2

genes, biofilm regulatory genes, and flagella.

DISCUSSION

Salmonella gains access to the systemic sites by traversing the gut intestinal barrier via utilizing

several routes. Its primary site of entry is the distal ileum wherein it induces the SPI-1 Type

Three Secretion System (T3SS) effector proteins to facilitate its uptake. After its initial entry

into the enterocytes and M-cells overlying the Peyer's patches, *Salmonella* colonizes the

draining MLN and eventually disseminates to the liver and spleen via the circulating

lymphatics and the bloodstream (40). *Salmonella* uptake by M cells facilitates its stealthy entry via avoidance of the mucus barrier. Additionally, the presence of flagella allows them to propel through the viscous mucus and activate the immune responses(16). The initial induction of pro-inflammatory responses during activation of cell-death pathways promotes systemic dissemination (40) (41, 42). The onset of systemic dissemination coincides with GVB disruption and loss of tight-junction protein expression (1). Several reports have suggested the protective role of gut commensals in maintaining gut epithelial health against food-borne pathogen insults. *E. coli* provide colonization resistance against STM in the presence of a specific microbial consortium by limiting galactitol wherein *Lachnospiraceae* consume the free sugar, conferring colonization resistance (43). Similarly, the commensal *Klebsiella michiganensis* retention or transmission provided colonization resistance to Enterobacteriaceae by exhibiting nutrient competition (44). In this study, we report the broad-spectrum protective role of the gut commensal of bacteroidales order *O. splanchnicus* against *Salmonella* and *Listeria* colonization, pathogenesis, and dissemination by inhibiting GVB damage, angiogenesis, inflammatory responses, and tight junction disruption in mice. Additionally*,* not only OS but its culture supernatant could exert its inhibitory effect on *Salmonella* invasion and intracellular proliferation in intestinal human epithelial cells. Moreover, the OS and its supernatant could inhibit STM biofilm formation by regulating flagella formation. This STM biofilm inhibitory effect was however lost upon co-culture with paraformaldehyde-fixed-OS indicating the presumable role of certain inhibitory microbial products secreted by metabolically live OS. Further, our studies identified OS-produced bacteriocins responsible for restricting intracellular replication of *Salmonella* and *Listeria* in intestinal epithelial cells by regulating bacterial virulence arsenals such as biofilm regulatory genes, SPI-1, SPI-2 genes, and flagellar machinery. However, the presence of flagella on *Salmonella* post bacteriocin treatment hints towards the possibility of other secreted accessory molecules within the OS

supernatant and warrants future research. Due to the proteinaceous nature of the bacteriocin,

oral administration experiments in mice were aborted. This study provides evidence about

inter-species regulation of gut colonization in maintaining gut homeostasis and protecting

against several invading pathogens. Moreover, the long-term protective ability conferred by

OS during pre-or post-infection colonization to mice against broad-spectrum foodborne

pathogens such as *Salmonella* and *Listeria* highlights its therapeutic potential as a prebiotic for

prophylaxis or post-infection treatment option, paving the way for future therapeutic

interventions to thwart food-borne infections.

MATERIALS AND METHODS

Bacterial Strains and Culture Conditions

Salmonella enterica serovar Typhimurium (STM) strain ATCC 14028S, STM ATCC 14028S

constitutively expressing mCherry (pFPV25.1-mCherry-Amp^R, RFP), laboratory stock of

Enterococcus faecalis (Ef), *Shigella flexneri*, *Listeria monocytogenes* expressing mCherry and

Odoribacter splanchnicus (OS) strain NCTC 10825 [1651/6] from ATCC were used in this

study. 4% paraformaldehyde-fixed-OS was used as killed bacteria control. The STM live

bacterial strain was grown overnight in LB broth at 37 ^{0}C under 160 rpm shaking condition.

OS was cultured in anaerobic condition in tryptic soy broth at 37ºC for 48h after its revival

from glycerol stock.

Preparation of Cell-free Bacterial Culture Supernatant

419 The overnight grown OS culture in tryptic soy broth was centrifuged at 10,000rpm for 10 min

420 and the supernatant was harvested. The collected supernatant was further subjected to another

421 two rounds to centrifugation to remove any residual cells. The supernatant was subsequently

422 filtered using a 0.2 µm filter and stored at − 20˚C.

Animal Experiment

The animal experiments were conducted in accordance with the approved guidelines of the institutional animal ethics committee at the Indian Institute of Science, Bangalore, India (Registration No: 48/GO/ReRcBiBt-S/Re-L/99/CPCSEA). All procedures involving the use of animals were performed according to the Institutional Animal Ethics Committee (IAEC)-approved protocol (CAF/Ethics/853/2021). 6–8 weeks old adult C57BL/6 male mice were used in this study. After one day of OS colonization (10^5 CFU/mice orally), the mice were infected with 10^6 (organ burden) CFU or 10^7 CFU (for assessing survival) of *S*. Typhimurium, 10^8 (organ burden) or 10^9 (survival) CFU of *L. monocytogenes* or 10^7 (organ burden) or 10^8 (survival) CFU of *S. flexneri* orally. The colonization of *O. splanchnicus* was confirmed via RT-qPCR mediated quantitation of OS 16S rRNA from the mice fecal and ileal samples. Infected mice were subjected to intraperitoneal injection of either 1 mg/kg body weight of SIRT1 inhibitor EX-527 (Sigma-Aldrich), or SIRT3 inhibitor 3TYP (Selleck Chemical) or vehicle control alone on every alternate day. For post-infection OS administration studies, OS was orally administered (10^5 CFU/ mice) post 1 or 3dpi. 5-day post-infection, mice were euthanized, dissected out, and blood and organs were harvested. The bacterial organ load was enumerated by plating the tissue homogenates on *Salmonella-Shigella* (SS) agar plates for *Salmonella* or *Shigella* or LB plates containing 50µg/ml of Ampicillin for LM. The harvested tissues and the fecal samples were collected in protein lysis buffer and TRIzol (Takara) for protein and RNA extraction respectively and were stored at -80ºC. For calculating percentage survival, the infected mice cohorts were monitored till fatality and fatalities were reported.

ELISA

Estimation of cytokines in mice serum was performed according to the manufacturer's instructions. Briefly, 96-well ELISA plates (BD Bioscience) were coated overnight with

capture antibody at 4°C. Following day, plates were washed with 0.1% Tween-20 containing

PBS and blocked with 10% FBS for 1 h. Following blocking, wells were washed and incubated

with 50µl of serum samples in duplicates for 2 h at room temperature. Subsequently, plates

were washed and incubated with detection antibody and enzyme reagent for 1 h at room

temperature (BD Bioscience). TMB (Sigma) was used as a substrate and reactions were stopped

with 2 N H_2SO_4. For the estimation of IL-1β, pre-coated ELISA (SARD Biosciences) plates

were used, and ELISA was performed as per manufacturer's protocol.

Absorbance was measured at 450 nm wavelength in Tecan Plate reader and the concentration

of cytokines were interpolated from a standard curve.

Quantitative Real Time PCR

Total RNA was isolated from tissues harvested in TRIzol (Takara) as per manufacturer's

protocol. RNA was quantified using Nano Drop (Thermo-Fischer scientific). Integrity of

isolated RNA was detected by performing 2% agarose gel electrophoresis. 3 µg of total isolated

RNA was subjected to DNaseI (Thermo Fischer Scientific) treatment at 37°C for 1 hr followed

by heat inactivation at 75°C in the presence of 5mM concentration of EDTA for 10 mins. The

mRNA was reverse transcribed to cDNA using oligo $(dT)_{18}$ primer, buffer, dNTPs and reverse

transcriptase (Takara) as per manufacturer's protocol. The expression profile of target gene was

evaluated using specific primers (Table-S2) by using SYBR green RT-PCR master mix

(Takara) in Applied Biosystems (Quantstudio 5) or BioRad Real time PCR instrument. β-actin

was used as an internal control for mammalian genes and for bacterial genes 16S rRNA was

used. All the reaction was setup in 384 well plate with three replicates for each sample.

Western blotting

The harvested tissue samples were lysed in ice-cold RIPA buffer (50 mM Tris-Cl, pH 8.0, 150

mM NaCl, 5mM EDTA, 1% NP-40, 1 % Sodium deoyxycholate, 1% SDS) containing protease

inhibitor cocktail (Roche). Total protein was estimated using Bradford reagent (BioRad).

Protein samples were denatured in **Laemmli SDS loading buffer** at 95ºC for 10 min and then

subjected to separation on 12% SDS-PAGE gels followed by transfer onto activated 0.45µm

PVDF membrane. Finally, the membrane was blocked with 5% skimmed milk in TBST (Tris

Buffered Saline containing 0.1% Tween-20) and probed with appropriate primary (Table-S3)

and secondary HRP-conjugated antibodies. The blots were developed using ECL substrate

(Bio-Rad) using a Chemidoc (BioRad). All densitometric analysis was performed using

ImageJ software.

Immunohistochemistry

The harvested tissues were fixed in 4% paraformaldehyde, embedded in paraffin and 5 µm

thick paraffin sections were collected on poly-lysine coated glass slides. The sectioned tissue

samples were deparaffinized using standard protocol. Primary antibody staining of the

deparaffinized sections was performed with specific primary antibody in the presence of a

permeabilizing agent, 0.01% saponin (Sigma) dissolved in 2.5% BSA containing PBS at 4°C

for overnight. Following wash with PBS, sections were stained with appropriate secondary

antibody tagged with fluorochrome for 1 hr at RT. This was followed by DAPI staining and

mounting. For in vivo estimation of biofilm, deparaffinized tissue samples were stained with

Calcofluor White (Sigma) stain for 20 min at RT. All immunofluorescence images were

obtained using Zeiss multi-photon 880 microscope and were analyzed using ZEN black 2012

software.

Haematoxylin and Eosin Staining

The harvested tissues were collected and fixed using 3.5% paraformaldehyde. The fixed liver

was then dehydrated using a gradually increasing concentration of ethanol and embedded in

paraffin. 5μm sections were collected on coated plates. Sections were further rehydrated and then stained with hematoxylin and eosin. Images were collected in a Leica microscope.

Cell Culture

Human colorectal adenocarcinoma cell line Caco-2 were maintained in Dulbecco's Modified Eagle's Media (DMEM) (Lonza) and supplemented with 10% FCS (Fetal calf serum, Gibco) and 1% non-essential amino acid solution (Sigma Aldrich) at 37ºC temperature in the presence of 5% CO_2 in humidified incubator. Prior to each experiment, cells were seeded into 24 well or 6 well plate as per requirement at a confluency of 60%.

Intracellular proliferation or gentamicin protection assay

Caco-2 epithelial were infected with 3 to 4 h old log phase *S*. Typhimurium (STM) bacterial culture, STM with *O. splanchnicus*, STM in presence of *O. splanchnicus* supernatant and STM along with PFA-fixed OS at an MOI of 10. For attachment of bacteria to the cells, tissue culture plates were subjected to centrifugation at 600xg for 5 min and incubated at 37 ^{0}C humified incubator with 5% CO_2 for 25 min. Following infection of the cells with STM at an MOI of 10, cells were treated with DMEM (Sigma) + 10% FBS (Gibco) containing 100 μg/ml gentamicin for 1 hr. Subsequently, the gentamicin concentration was reduced to 25 μg/ml and maintained until the specified time point. Post 2hr and 16hr post-infection, cells were lysed in 0.1% triton-X-100. Lysed cells were serially diluted and plated on *Salmonella-Shigella* (SS) agar to obtain colony-forming units (cfu). Percent invasion was determined using the following formula:

Percent invasion= [CFU at 2 h]/ [CFU of pre-inoculum] X 100

Fold proliferation was calculated as cfu at 16hr divided by cfu at 2hr.

Fold Proliferation= [CFU at 16h] / [CFU at 2h]

Immunofluorescence confocal microscopic studies

At the specified time points post infection with mCherry-tagged STM, or GFP-tagged OS or

with both mCherry-STM and GFP-OS, cells were fixed with 3.5% paraformaldehyde for 15

min. Primary antibody staining was performed with specific primary antibody in the presence

of a permeabilizing agent, 0.01% saponin (Sigma) dissolved in 2.5% BSA containing PBS at

4°C for overnight or for 6hr at room temperature (RT). Following this, cells were washed with

PBS stained with appropriate secondary antibody tagged with fluorochrome for 1 hr at RT. This

was followed by DAPI staining and mounting of the coverslip onto a clean glass slide using

the mounting media containing the anti-fade agent. The coverslip sides were sealed with a

transparent nail paint. All immunofluorescence images were obtained using Zeiss Multiphoton

880 and were analyzed using ZEN black 2012 software.

Growth Curve analysis

For *in vitro* growth curve experiments a single colony of STM and OS was inoculated in 5mL

of LB broth singly or in co-culture and grown overnight at 37ºC in anaerobic jar. In another

set, STM was inoculated in 5mL of LB in presence of OS supernatant. Overnight-grown

stationary phase bacteria were sub-cultured at a ratio of 1: 100 in freshly prepared LB in

presence or absence of OS supernatant and kept at 37ºC. At different time intervals, aliquots

were taken for measurement of $[OD]600nm$ by TECAN 96 well microplate reader and were

plated for CFU enumeration.

Growth conditions for biofilm formation

538   LB medium devoid of NaCl (i.e. 1% casein enzyme hydrolysate (Hi-Media, India) and 0.5%

539   yeast extract (Hi-Media, India), devoid of NaCl) was used as biofilm media. Overnight grown

540   culture were subcultured in 2 ml biofilm medium at the dilution of 1:100 in a flat-bottom 24-

541   well polystyrene plate (Tarsons, India) and incubated at 28˚C for 72 h without shaking.

Bacterial Biofilm inoculation with culture supernatant

To test the effect OS culture supernatant on biofilm formation, 20 µl of STM WT overnight culture and indicated volume (1% v/v and/or as indicated in the figures) of the OS culture supernatants was added to 2 ml of biofilm media,and incubated under biofilm inducing condition. For co-culture experiments, the overnight cultures of the STM and OS or STM and PFA-fixed OS were added in a 1:1 ratio to 2 ml of biofilm media and incubated under biofilm-inducing conditions.

Crystal Violet Staining

Post 72 h of biofilm induction, the media was discarded, and the plates were thoroughly washed in water. The plates were dried, and 2 ml of 1% Crystal Violet (CV) was added to each well and incubated for 15 min. Post 15 min incubation, the CV stain was removed, and the plate was thoroughly rinsed in water to remove excess stain. The stained biofilm was destained with 70% ethanol, and the intensity of the colour of the destained solution was quantified at OD_{595} in the Tecan plate reader (Infinite Pro 200).

Glass bead assay

1 mm glass beads were added onto the pellicle to assess the strength of the pellicle at the air-liquid interface. The initial weight of the glass beads was noted. The glass beads were added until the pellicle collapsed, and the final weight was noted down and plotted.

Gut Permeability Assay

The infected mice cohorts were orally administered with FITC Dextran (25mg/kg,FD-4-100MG, Sigma) 4h before euthanization. The blood plasma was collected post-euthanization by cardiac puncture followed by centrifugation. The blood plasma was evaluated for mean fluorescence via plate reader (TECAN) and flow cytometry(CytoFlex, Beckman).

MTT Assay

Post-16h treatment of pre-seeded Caco-2 cells with 10µg/ml of bacteriocin, MTT staining solution (0.5mg/ml) was added to each well followed by incubation at 37ºC CO_2 incubator for 4h. 50µl of DMSO was added per well for solubilizing the formazan crystals. Absorbance was measured at 570nm using a plate reader (TECAN).

Transmission electron microscopy

Overnight STM or LM-grown cultures (in the presence or absence of OS supernatant or bacteriocin) were subcultured into fresh media (1:10) ratio and incubated at 37ºC for 6 hours under shaking conditions. These samples were centrifugated at 1000rpm for 10 min to obtain the bacterial pellet. The bacterial pellet was subjected to PBS wash twice via centrifugation at 1000 rpm for 10 min.5 µl of the finally resuspended cell suspension was added on a glow-discharged copper-grid, air dried, stained with 1% uranyl acetate for 30 s, and visualized under a transmission electron microscope (JEM-100 CX II; JEOL).

Supernatant Treatment

The supernatant was treated with 7.5 mM concentrations of EDTA and incubated at 65 °C for 1 h. To 200 µl of supernatant, 10 µl of proteinase-K (NEB, stock 20 mg/ml) was added and incubated at 37 °C for 1 h. The proteinase was inactivated with 0.5 mM PMSF and incubated at 28 °C for 1 h. The supernatant was heated to 37˚C (60 min), 65˚C (15 min) or 95˚C (15 min) for the indicated time and immediately frozen at -20˚C. Post-treatment, the supernatants were used to inoculate the biofilm to test the nature of the biofilm inhibitory molecules.

Supernatant concentration and separation based on molecular weight

4 ml of the supernatants were transferred to the Amicon ultra filter device (Amicon® Ultra-4 centrifugal filter device, 3k MWCO, UFC800324; Amicon® Ultra-4 Centrifugal filter Unit, 10k MWCO, UFC801024) and centrifuged at 4000 g for 30 min in a swing bucket rotor. The concentrated solute from the bottom of the filter device and the flowthrough were collected by a pipette and subjected to biofilm inoculation.

Size Exclusion Chromatography

The bacterial supernatant was concentrated to 0.5ml using 3k MWCO-Amicon-Ultra-4-centrifugal unit and loaded onto the Superdex 200pg size exclusion column (XK16/60, 124mL; Cytiva, USA), pre-equilibrated Tris-Cl buffer (pH-8) containing 100mM NaCl and 20mM Tris-Cl. The sample was eluted with the same buffer at a flow rate of 0.8mL/min, and 1mL fractions were collected. The fractions corresponding to the elution peaks were analyzed for biofilm inhibition activity and subjected to mass spectrometry.

LC/MS analysis of the supernatant fractions

The concentrated gel filtration fractions of supernatants were subjected to in-solution trypsin digestion for LC Q-TOF MS analysis. Briefly, 100 μg of total protein was resuspended in 100 μl of 7-8M of urea buffer. To this, 5 μl 200mM DTT was added and incubated at room temperature for 1 hr, followed by alkylation with 20µl of 200 mM iodoacetamide and incubation at room temperature for 1h in the dark. This was followed by the addition of 1mM of Calcium chloride and incubation at RT for 10 min. For protein digestion, trypsin solution was added to the sample to a final trypsin to protein ratio of 1:30 (w/w) and incubated at 37˚C for 16 h, with frequent shaking. The reaction was stopped by adding 0.1% formic acid to adjust the pH to 3-4. The digested protein samples were desalted using a reverse phase C-18 column

(Acclaim prep EASY-Spray (C18-RP)) using mobile phase solvent A-2% acetonitrile and

solvent B- 80% acetonitrile. The sample injection volume was 10µl. All the samples were

analyzed using Thermo Scientific SII for Xcalibur method. The MS scan was acquired using

Orbitrap detector with 60000 resolution in the positive polarity and scan range (*m/z*) 350-2000.

The ion source type was NSI with static voltage; positive ion (V)- 2200V and negative ion (V)-

600 along with temperature settings at 275ºC. This was followed by ddMS^2 IT HCD with 30%

fragmentation HCD collision energy using ion trap detector. The raw data was analyzed using

Proteome Discoverer 2.5 against the *O. splanchnicus* proteome available in UniProt (Proteome

ID: UP000006657).

Purification of bacteriocin

O. splanchnicus culture supernatant was concentrated within the range of >3kDa and <10kDa

using the Amicon Ultra-4 Centrifugal filter units of 10MWCO and 3MWCO. The concentrated

cell-free supernatant containing the bacteriocin (~7kDa) was precipitated by ammonium

sulphate (80% to 98%). The precipitate was collected by centrifugation at 12000rpm for 30

min and dissolved in 1X PBS. The dissolved precipitate was again filtrated using the 3KDa

cutoff Amicon centrifugal filter units. This concentrated fraction was further subjected to Fast

Performance Liquid Chromatography using Superdex 75 Increase 10/300 GL- 24ml volume

(GE Healthcare) column. The column was equilibrated with 1X Phosphate Buffered Saline

buffer and 0.5ml of concentrated sample (23.6 mg/ml) was injected into the column. Based on

the increase in Absorbance units at 280nm from baseline, 0.3ml fractions were collected at a

retention volume of 13.3ml to 21.8ml (Fig. S9). The fractions were analyzed by 20% SDS-

PAGE electrophoresis and stained with Coomassie staining solution.

In vitro **swim motility assay**

634 2 μl of treated or untreated bacterial samples were spotted on the 0.3% agar (LB-STM, BHI-

635 LM) plates supplemented with 0.5% yeast extract, 1% casein enzyme hydrolysate, 0.5% NaCl,

636 and 0.5% glucose. The plates were incubated at 37°C and images were taken every 6 h (STM)

637 or 24h (LM) using the camera of Biorad illuminator.

Statistical analysis

639 Data were analyzed and graphed using the GraphPad Prism 8 software (San Diego, CA).

640 Statistical significance was determined by Student's t-test or Two way ANOVA and Bonferroni

641 post-t-test to obtain p values. Adjusted p-values below 0.05 are considered statistically

642 significant. The results are expressed as mean ± SD or SEM of three independent experiments.

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MAIN FIGURES

Figure 1. *Odoribacter splanchnicus* **colonization protects mice from** *Salmonella*

colonization and dissemination

A-Schematic representation showing animal experimental strategy of infection.

B-OS colonization validation via qPCR quantification of *O. splanchnicus* 16S rRNA from fecal

samples of mice.

C-Survival percentage of non-colonized-STM-infected mice versus OS-colonized-STM-

infected mice cohorts. Data is represantative of N=2, n=5.

D-Percent weight loss exhibited by OS colonized and non-colonized mice upon STM infection

at 5th day post infection. Data is representative of N=3,n=5.

E-G- STM organ loads in MLN (E), liver (F) and blood (G) at 5th post-infection in OS colonized

and non-colonized mice cohorts in untreated, and SIRT1 (EX-527) or SIRT3 (3TYP) inhibitor

treatment. Data is representative of N=3, n=3. Mann Whitney Test was performed to obtain p-

values.

Figure 2. *O. splanchnicus* reduces *Salmonella* induced inflammation and intestinal

damage

A-Estimation of serum IL-6 production via ELISA upon STM infection in OS non-colonized

and colonized mice cohorts in presence or absence of SIRT1 or SIRT3 inhibitor treatment.

Unpaired t-test is performed to obtain the p-values. Data is representative of N=2, n=5.

B-Quantitative gene expression data of several antimicrobial peptides (AMPs) precursor genes

in uninfected and STM-infected mice intestinal tissues harvested at 5th dpi from OS colonized

or non-colonized cohorts. Unpaired t-test is performed to obtain the p-values. Data is representative of N=2, n=5.

C- Immunostaining representing Muc2 protein expression in intestinal tissues of OS non-colonized and colonized mice cohorts upon STM infection.

D-Haematoxylin-eosin-stained intestinal sections harvested from STM-infected OS non-colonized or colonized mice cohorts at 5th dpi. 0-represents normal pathology, 1- represents-mild pathology, 2-3-represents severe pathology.

Fig 3. *Odoribacter splanchnicus* gut colonization prevents *Salmonella*-induced GVB damage by promoting tight junction gene expression.

A-Quantitative gene expression studies of tight junction protein complexes (*Tjp1*, *Claudin5*) and PV-1 gene (marker of GVB damage) in STM-infected mice intestinal tissue in presence or absence of OS gut colonization.

B-Immunoblotting depicting expression of PV-1 and CD34 protein in intestinal tissues of OS

non-colonized and colonized mice cohorts upon STM infection.

C-Immunohistochemistry representing CD34 (endothelial marker of vasculature) protein

expression in intestinal tissues of OS non-colonized and colonized mice cohorts upon STM

infection.

D-Quantitative representation of the Median Fluorescence Intensity (MFI) of CD34 protein

expression in intestinal immunohistochemical sections shown in (C). Data is representative of

N=2,n>50. Un-paired t-test was used to obtain the p-values.

E- Estimation of FITC fluorescence in mice plasma via gut permeability assay in infected mice

cohorts upon oral administration of FITC Dextran (25mg/kg) 4h before euthanization at 5th day

post *Salmonella* infection.

Fig 4. *O. splanchnicus* **exhibits its inhibitory effect on** *Salmonella* **biofilm formation and**

inhibits invasion and intracellular replication in human intestinal cells

A-Percent invasion of STM in human Caco-2 human intestinal cells in presence or absence of

live OS, OS culture supernatant, PFA-fixed-OS. Data is representative of N=3, n=3. Unpaired

t-test is used to obtain the p-values.

B-Intracellular proliferation of STM Caco-2 cells in presence or absence of live OS, OS culture

supernatant, PFA-fixed-OS. Data is representative of N=3, n=3. Unpaired t-test is used to

obtain the p-values.

C- Representative immunofluorescence images depicting STM invasion into Caco-2 cells in presence or absence of OS at 2hr post-infection. Data is representative of N=3, n=100. Unpaired t-test is used to obtain the p-values.

D- Representative immunofluorescence images depicting STM invasion into Caco-2 cells in presence or absence of OS at 2hr post-infection. Data is representative of N=3, n=100. Unpaired t-test is used to obtain the p-values.

E-Percent invasion of STY in Caco-2 human cells in presence or absence of OS, OS culture supernatant, PFA-fixed OS. Data is representative of N=3, n=3. Unpaired t-test is used to obtain the p-values.

F-Intracellular proliferation of STY in Caco-2 cells in presence or absence of live OS, OS culture supernatant, PFA-fixed-OS. Data is representative of N=3, n=3. Unpaired t-test is used to obtain the p-values.

G- Biofilm assay showing the biofilm formation ability of STM-WT or STM-WT grown in OS supernatant after 72h of inoculation in presence or absence of live OS, OS culture supernatant, PFA-fixed-OS. Data is representative of N=4, n=4. Unpaired t-test is used to obtain the p-values. Image of the STM biofilm formed plate in the above stated conditions has been shown.

H-M- RT-qPCR mediated gene expression of STM biofilm regulatory genes (*csgD*) (H), *csgA* (I), *bapA* (J), along with flagellar gene (*fliC*) (K), SPI-1 (*sipC*) (L) and SPI-2 (*pipB2*) (M) genes in presence or absence of live OS or its spent media or PFA-fixed-OS. Data is representative of N=2, n=5. Unpaired Student's t-test was used to obtain the p-values.

N- Transmission Electron Microscopic (TEM) images of STM in presence or absence of *Salmonella* WT supernatant, OS and its supernatant to assess the possession of flagella.

Fig 5. Unlike *O. splanchnicus, E. faecalis* fail to exert any inhibitory effect on *Salmonella in vitro* biofilm formation or *Salmonella* invasion and intracellular replication in human intestinal cells

851 A-Percent invasion of STM in Caco-2 cells in presence or absence of live *E. faecalis* (Ef), Ef culture supernatant, PFA-fixed-Ef. Data is representative of N=3, n=3. Unpaired t-test is used to obtain the p-values.

854 B-Intracellular proliferation of STM in Caco-2 cells in presence or absence of live Ef, Ef culture supernatant, PFA-fixed-Ef. Data is representative of N=3, n=3. Unpaired t-test is used to obtain the p-values.

857 C. Biofilm assay depicting the biofilm formation ability of STM-WT or STM-WT grown in Ef supernatant after 72h of inoculation in presence or absence of live Ef, Ef or STM culture supernatant, PFA-fixed. Crystal-violet staining was performed to assess the biofilm formation on the solid-liquid interphase and the [OD]$_{595}$ absorbance reading was recorded. Data is representative of N=4, n=4. Unpaired t-test is used to obtain the p-values.

Fig 6. *Odoribacter splanchnicus* exerts its protective role against *Listeria monocytogenes* but not against *Shigella flexneri* in mice

A- Survival percentage of non-colonized LM-infected mice versus OS-colonized *Listeria*-infected mice cohorts. Data is represantative of N=2, n=8.

B-Percent weight loss exhibited by OS colonized and non-colonized mice upon LM infection at 5th dpi. Data is representative of N=2,n=8.

C-F- LM organ loads in Mesenteric lymph Node (MLN) (C), liver (D), spleen (E) and blood (F) at 5th dp. in OS colonized and non-colonized mice cohorts Data is representative of N=2, n=8. Mann Whitney Test was performed to obtain p-values.

G- Survival percentage of non-colonized *Shigella*-infected mice versus OS-SF-infected mice cohorts. Data is represantative of N=2, n=8.

H-Percent weight loss exhibited by OS colonized and non-colonized mice upon *S. flexneri* infection at 5th dpi. Data is representative of N=2,n=8.

878 I-L- *S. flexneri* organ loads in Mesenteric lymph Node (MLN) (I), liver (J), spleen (K) and

879 blood (L) at 5ᵗʰ dpi in OS colonized and non-colonized mice cohorts. Data is representative of

880 N=2, n=8. Mann Whitney Test was performed to obtain p-values.

Fig. 7 *O. splanchnicus* **even confers protection against foodborne pathogens when administered post-pathogenic challenge**

A- Schematic representation showing the animal experimental strategy of infection.

B- Percent weight loss exhibited by OS colonized (1-dpi) and non-colonized mice upon LM infection at 5th dpi. Data is representative of N=2,n=8.

C-F- LM organ loads in MLN (C), liver (D), spleen (E) and blood (F) at 5th dpi upon OS administration at 1-dpi. Data is representative of N=2, n=8. Mann Whitney Test was performed to obtain p-values.

G- Percent weight loss exhibited by OS colonized (1-dpi) and non-colonized mice upon STM infection at 5th dpi. Data is representative of N=2,n=8.

H-K- STMorgan loads in MLN (H), liver (I), spleen (J) and blood (K) at 5th-dpi upon OS administration at 1-dpi. Data is representative of N=2, n=8. Mann-Whitney Test was performed to obtain p-values.

L- Survival percentage of non-colonized LM-infected mice versus OS-colonized (1 dpi or 3 dpi) *Listeria*-infected mice cohorts. Data is represantative of N=2, n=8.

M- Percent weight loss exhibited by OS colonized (3-dpi) and non-colonized mice upon LM infection at 5th-day post-infection. Data is representative of N=2,n=8.

N-Q- LM organ loads in MLN (N), liver (O), spleen (P) and blood (Q) at 5th-dpi upon OS administration at 3-dpi. Data is representative of N=2, n=8. Mann-Whitney Test was performed to obtain p-values.

R- Survival percentage of non-colonized *Salmonella*-infected mice versus OS-colonized (1 dpi or 3 dpi) *Listeria*-infected mice cohorts. Data is represantative of N=2, n=8.

S- Percent weight loss exhibited by OS-colonized (3-dpi) and non-colonized mice upon STM infection at 5th-day post-infection. Data is representative of N=2,n=8.

T-W-STM organ loads in MLN (T), liver (U), spleen (V), and blood (W) at 5th-dpi upon OS

administration at 3-dpi. Data is representative of N=2, n=8. Mann-Whitney Test was performed

to obtain p-values.

X- Haematoxylin-eosin-stained intestinal sections harvested at 5th-dpi from Uninfected (UI) or

Salmonella or *Listeria*-infected mice cohorts having non-administered or administered OS (at

1st or 3rd dpi). 0-represents normal pathology, 1- represents- mild pathology, 2-3-represents

severe pathology.

Y- Representative Calcofluor white (CW) staining of STM-infected mice spleen tissues at 5th-

day post-infection with or without OS administration at 1st or 3rd dpi

Z- Representative CW staining of LM-infected mice spleen tissues at 5th-dpi with or without

OS administration at 1st or 3rd dpi.

I   STM   STM+ *O. splanchnicus* supernatant   STM+ *O. splanchnicus* bacteriocin   **J**   LM   LM+ *O. splanchnicus* supernatant   LM+ *O. splanchnicus* bacteriocin

Fig. 8 *O. splanchnicus* bacteriocin restricts both *Salmonella* and *Listeria* pathogenesis

A- Percent invasion and intracellular fold proliferation of STM in Caco-2 cells under OS bacteriocin (10µg/ml) treatment. Data is representative of N=2, n=3. Unpaired t-test is used to obtain the p-values.

B- Percent invasion and intracellular fold proliferation of LM in Caco-2 cells under OS bacteriocin (10µg/ml) treatment. Data is representative of N=2, n=3. Unpaired t-test is used to obtain the p-values.

C- Percent invasion and intracellular fold proliferation of STY in Caco-2 cells under OS bacteriocin (10µg/ml) treatment. Data is representative of N=2, n=3. Unpaired t-test is used to obtain the p-values.

D-MTT assay evaluating the viability of Caco-2 cells upon 16h of mock or bacteriocin (10µg/ml) treatment. Data is representative of N=2, n=4.

E-F-TEM images of STM under untreated (D) or bacteriocin (10µg/ml) (E) treatment to assess the possession of flagella.

G-H- TEM images of LM under untreated (D) or bacteriocin (10µg/ml) (E) treatment to assess the possession of flagella.

I- Swimming motility assay of STM under OS supernatant or bacteriocin treatment.

J- Swimming motility assay of LM under OS supernatant or bacteriocin treatment.

List of Supplementary Materials

Fig. S1 to S9

Table S1 to S3

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All data are available in the main text or the supplementary materials.

SUPPLEMENTARY FIGURES

Fig S1. *O. splanchnicus* **gut colonization prevents** *Salmonella* **dissemination by restricting intestinal GVB damage and angiogenesis**

A-Immunohistochemistry demonstrating PV-1 protein expression in intestinal tissues of OS non-colonized and colonized mice cohorts upon STM infection.

B-C Immunohistochemical sections representing CD34 (an endothelial marker of vasculature) protein expression in intestinal (B) and liver (C) tissues of OS non-colonized and colonized mice cohorts upon STM infection. Data is representative of N=2, n=6 (tissue samples). Un-paired t-test was used to obtain the p-values.

D- qPCR mediated gene expression of pro-angiogenic marker *Vegf* in STM-infected mice liver tissues in presence or absence of OS pre-colonization. Data is representative of N=2, n=5. Un-paired t-test was used to obtain the p-values.

E- qPCR mediated gene expression of *Axin2* in STM-infected mice intestinal tissues in the presence or absence of OS pre-colonization. Data is representative of N=2, n=5. Un-paired t-test was used to obtain the p-values.

Fig. S2 In vitro growth kinetics of *Salmonella* Typhimurium in presence of live or dead *O. splanchnicus* or its culture supernatant

A-B Growth kinetics of STM-WT, OS, STM-WT in presence of OS, STM-WT in presence of STM or OS culture supernatant and STM-WT in presence of PFA-fixed OS in LB under anaerobic growth conditions.

C-D- Growth kinetics of STM-WT, OS, STM-WT in presence of OS, STM-WT in presence of STM or OS culture supernatant, STM-WT supernatant, and STM-WT in presence of PFA-fixed OS in LB under aerobic growth conditions.

Fig. S3. *Odoribacter splanchnicus* **gut colonization protects mice against** *Listeria monocytogenes* **pathogenesis similar to** *Salmonella* **infection**

A-Haematoxylin-eosin-stained intestinal sections harvested from OS non-colonized or colonized mice cohorts at 5th-day-post-LM-infection. 0-represents normal pathology, 1-represents- mild pathology, 2-3-represents severe pathology.

B-Quantitative gene expression data of several antimicrobial peptides (AMPs) precursor genes in LM-infected mice intestinal tissues harvested at 5th day post-infection from OS colonized or non-colonized cohorts. Unpaired Student's t-test is performed to obtain the p-values. Data is representative of N=2, n=5.

C- Haematoxylin-eosin-stained intestinal sections harvested from OS non-colonized or colonized mice cohorts at 5th-day-post-SF-infection. 0-represents normal pathology, 1-represents- mild pathology, 2-3-represents severe pathology.

D- Quantitative gene expression data of several antimicrobial peptides (AMPs) precursor genes in SF-infected mice intestinal tissues harvested at 5th-dpi from O.S colonized or non-colonized cohorts. Unpaired Student's t-test is performed to obtain the p-values. Data is representative of N=2, n=5.

E- Quantitative gene expression study of tight junction protein genes *Tjp1*, *Claudin5*, GVB damage marker *PV-1*, angiogenesis marker-*Vegf*, and *Axin2* in LM- infected mice intestinal tissues harvested at 5th-dpi from OS colonized or non-colonized cohorts. Unpaired Student's t-test is performed to obtain the p-values. Data is representative of N=2, n=5.

F- Quantitative gene expression study of tight junction protein genes *Tjp1*, *Claudin5*, GVB damage marker *PV-1*, angiogenesis marker-*Vegf*, and *Axin2* in SF-infected mice intestinal tissues harvested at 5th-dpi from OS-colonized or non-colonized cohorts. Unpaired Student's t-test is performed to obtain the p-values. Data is representative of N=2, n=5.

G-H- Intestinal Immunohistochemical sections representing PV-1 protein expression (G) and Muc2 (H) in intestinal tissues of OS-non-colonized and colonized mice cohorts upon LM infection.

I-J- Intestinal Immunohistochemical sections representing PV-1 protein expression (I) and Muc2 (J) in intestinal tissues of OS non-colonized and colonized mice cohorts upon SF infection.

K- Representative Calcofluor white (CW) staining of STM-infected mice spleen tissues at 5th-dpi.

L- Representative CW staining of LM-infected mice spleen tissues at 5th-dpi.

M- Transmission Electron Microscopic (TEM) images of LM in presence or absence of LM supernatant, OS and its supernatant to assess the possession of flagella.

N-O- Immunohistochemical sections representing CD34 (an endothelial marker of vasculature) protein expression in intestinal (N) and liver (O) tissues of OS non-colonized and colonized mice cohorts upon LM infection. Data is representative of N=2, n=3 (tissue samples). Un-paired t-test was used to obtain the p-values.

Fig. S4 *Odoribacter splanchnicus* **and its culture supernatant restricts** *Listeria monocytogenes* **pathogenesis in human intestinal cells**

A- Percent invasion of LM in Caco-2 cells in presence or absence of live OS, OS sup, PFA-fixed-OS. Data is representative of N=3, n=3. Unpaired t-test is used to obtain the p-values.

B-Intracellular fold proliferation of LM in Caco-2 cells in the presence or absence of live OS, OS sup, PFA-fixed-OS. Data is representative of N=3, n=3. Unpaired t-test is used to obtain the p-values.

Fig. S5 *Odoribacter splanchnicus* administration post-establishment of infection protected mice against *Listeria monocytogenes* and *Salmonella* Typhimurium

A-E-Quantitative gene expression data of several antimicrobial peptides (AMPs) precursor genes in LM and STM-infected mice intestinal tissues harvested at 5th dpi upon OS colonization at 1-dpi or 3-dpi. Unpaired Student's t-test is performed to obtain the p-values. Data is representative of N=2, n=5.

F-J- Quantitative gene expression study of tight junction protein genes *Tjp1*, *Claudin5*, GVB damage marker *PV-1*, angiogenesis marker-*Vegf,* and *Axin2* in LM and STM-infected mice intestinal tissues harvested at 5th-dpi upon OS colonization at 1-dpi or 3-dpi. Unpaired Student's t-test is performed to obtain the p-values. Data is representative of N=2, n=5.

K-L- Intestinal Immunohistochemical sections representing PV-1 protein expression in *Salmonella* (K) or *Listeria* (L)-infected mice harvested on 5th-dpi with or without OS administration at 1-dpi or 3-dpi.

M-N- Intestinal Immunohistochemical sections representing Muc2 protein expression in *Salmonella* (M) or *Listeria* (N)-infected mice harvested on the 5th dpi with or without OS administration at 1-dpi or 3-dpi.

O-P- Immunoblotting representing the protein expression profile of Muc2, ZO-1, and PV-1 in *Salmonella* (O) or *Listeria* (P)-infected mice intestinal tissue harvested on 5th-dpi with or without OS pre-or post-colonization (1-dpi or 3-dpi).

Q- Estimation of FITC fluorescence in mice plasma via gut permeability assay in *Salmonella* or *Listeria*-infected mice cohorts with or without OS pre-or post-colonization (1dpi or 3dpi) upon oral administration of FITC Dextran (25mg/kg) 4h before euthanization at 5th-dpi.

Fig. S6 *Odoribacter splanchnicus* alleviates *Salmonella*-induced intestinal inflammation in mice both during pre and post-treatment

A- Estimation of IL-1β in the ileal sections of UI or *Salmonell*a-infected mice tissues harvested at 5th dpi with or without OS pre-or post-colonization (1dpi).

Conditions or Parameters Tested	Range	Outcome
Concentration	0.4% v/v to 2% v/v	Biofilm inhibition and pellicle defect at all tested concentrations.
Temperature Sensitivity	37ºC, 65ºC and 95ºC	Biofilm inhibition and pellicle defect at all temperatures tested.
Size Cut off	<3kDa, >3kDa, <10kDa, >10kda	>3kDa-Biofilm inhibition <10kDa, >10kDa-Biofilm inhibition
Proteinase K treatment	Proteinase K treatment sensitive	Biofilm inhibition got reversed upon proteinase K treatment.
EDTA Treatment	EDTA treatment resistant	No effect on biofilm inhibition and pellicle defect upon EDTA.

Comparison of Gel Filtration fraction peaks of Protein Standard versus Sample

65kDa

➡️ Fractions showing inhibitory effect   🟢 Fractions exhibiting highest inhibitory activity upon titration

A82, A83, and A84- Fractions exhibiting highest inhibitory activity upon titration

Fig. S7 Characterization of the inhibitory active molecule within *O. splanchnicus* culture supernatant

A- Biofilm assay showing the biofilm formation ability of STM after 72h of inoculation in presence of OS culture supernatant upon EDTA treatment. Data is representative of N=4, n=4. Unpaired t-test is used to obtain the p-values.

B- Biofilm assay showing the biofilm formation ability of STM after 72h of inoculation in presence of OS culture supernatant at high temperature conditions. Data is representative of N=4, n=4. Unpaired t-test is used to obtain the p-values.

C- Biofilm assay showing the biofilm formation ability of STM after 72h of inoculation in presence of OS culture supernatant under Proteinase K treatment. Data is representative of N=4, n=4. Unpaired t-test is used to obtain the p-values.

D- Coomassie-stained SDS-PAGE demonstrating the size range of the differential protein bands present in OS supernatant as opposed to the tryptic soy broth media control.

E- Biofilm assay showing the biofilm formation ability of STM after 72h of inoculation in presence of varying concentration OS culture supernatant. Data is representative of N=4, n=4. Unpaired t-test is used to obtain the p-values.

F- Biofilm assay showing the biofilm formation ability of STM after 72h of inoculation in presence of OS culture supernatant fractions fractionated using 3kDa Amicon filter. Data is representative of N=4, n=4. Unpaired t-test is used to obtain the p-values.

G- Biofilm assay showing the biofilm formation ability of STM after 72h of inoculation in presence of OS culture supernatant fractions fractionated using 10kDa Amicon filter. Data is representative of N=4, n=4. Unpaired t-test is used to obtain the p-values.

H- Summary table demonstrating the conditions tested for active molecule characterization and their effective outcome.

I-Gel filtration chromatogram of protein standards and the OS culture supernatant. Red arrow shows the fractions with biofilm inhibitory activity. Green arrow depict the fraction with highest inhibitory activity.

J-K- Biofilm assay showing the biofilm formation ability of STM after 72h of inoculation in presence of OS supernatant gel filtration fractions. Data is representative of N=2, n=2. Unpaired t-test is used to obtain the p-values.

L- Biofilm assay showing the biofilm formation ability of STM after 72h of inoculation in presence of varying concentration of OS supernatant biofilm inhibitory gel filtration fractions. Data is representative of N=2, n=1. Unpaired t-test is used to obtain the p-values.

M-Schematic representation of the workflow of mass spectrometric analysis of the gel filtration fractions (A82, A83, A84 and Concentrated supernatant).

Fig. S8 *O. splanchnicus* bacteriocin attenuates *Salmonella* pathogenesis

A- Biofilm assay showing the biofilm formation ability of STM after 72h of inoculation under varying concentration of OS bacteriocin. Data is representative of N=2, n=2. Unpaired t-test is used to obtain the p-values.

B-G- RT-qPCR mediated gene expression of STM biofilm regulatory genes *csgD* (B), *csgA*(C), *bap* (D), along with flagellar gene *fliC*(E), SPI-1-*sipC* (F) and SPI-2-*pipB2*(G) in presence or absence of OS bacteriocin treatment. Data is representative of N=2, n=5. Unpaired Student's t-test was used to obtain the p-values.

Fig. S9 Purification procedure of *O. splanchnicus* produced bacteriocin via ammonium sulphate precipitation method followed by Fast Performance Liquid Chromatography

A-Size exclusion chromatogram of OS-produced bacteriocin fractions post 95-98% ammonium sulphate precipitation.

B- Coomassie-stained SDS PAGE showing the purified and concentrated OS bacteriocin

Table S1- Selected List of 11 proteins from Mass Spectrometric Analysis based on bibliographic search

Accession ID	Protein Name and	Referenced Function
A0A412WBT8	Immunity protein Imm5 domain-containing protein	Uncharacterized
F9Z8G5	Peptidase	Modulate biofilm formation and host defense.
R6F650	Carboxyl-terminal protease	Critical for post-translational processing and associated with regulation of gene expression, stress response, maintenance of cell envelope integrity and bacterial virulence.
A0A412TJE3	ATP-dependent Clp protease proteolytic subunit	Influence virulence and infectivity of pathogenic bacteria. ClpP enhance biofilm formation in *Enterococcus faecalis*.
F9Z604	Peptidase C39 bacteriocin processing	The cysteine peptidase cleave the double glycine leader peptides from the precursors of bacteriocin
A0A412TJV3	UDP-2,4-diacetamido-2,4, 6-trideoxy-beta-L-altropyranose hydrolase	It catalyzes the third step in the biosynthesis of pseudaminic acid. Pseudaminic acid is a sialic-like sugar that modifies flagellin.
A0A412WSU3	Bacteriocin	Antimicrobial peptides that can kill or inhibit closely related or unrelated bacterial strains.
A0A413I832	Serine Protease	Extracellular proteases regulate biofilm formation and growth of competitors.
A0A412W3J2	Glutamate dehydrogenase	Involved in bacterial response to stress adaptation and virulence.
A0A412WSS0	Bacteriocin	Antimicrobial peptides that can kill or inhibit closely related or unrelated bacterial strains.
F9ZBU9	Peptidylprolyl isomerase	Promote biofilm formation in *Mycobacterium tuberculosis*.

Table S2-List of primers

Primer Name	Sequence (5'---3')
O. splanchnicus 16S rRNA FP	ATGTAATGATGAGCACTCTAACGG
O. splanchnicus 16S rRNA RP	GGCTTTTGAGATTGGCATCC
PV-1 FP	CGAACTGGAGGCGCGTAA
PV-1 RP	GCAGCGACTTGGCCTTGAC
Claudin5 FP	CCCAGTTAAGGCACGGGTAG
Claudin5 RP	GGCACCGTCGGATCATAGAA
βactin FP	CAGCAAGCAGGAGTACGATG
βactin RP	GCAGCTCAGTAACAGTCCG
Vegf FP	GCAGCGACAAGGCAGACTAT

Vegf RP	AACCTCCTCAAACCGTTGGC
Tjp1 FP	TCTTGCAAAGTATCCCTTCTGT
Tjp1 RP	CAGAAATCGTGCTGATGTGCC
Reg3b FP	GCTTGTCAAGAGCTTCTGGATTTC
Reg3b RP	TGTCCCTTGTCCATGATGCTC
Reg3g FP	CGCTGAAGCTTCCTTCCTGT
Reg3g RP	GGGTTCATAGCCCAGTGTCG
Lysozyme C FP	GGTCTACAATCGTTGTGAGTTG
Lysozyme C RP	TGAGCTAAACACACCCAGTCA
CCL 20 FP	GCAACTACGACTGTTGCCTC
CCL20 RP	TTCTTGACTCTTAGGCTGAGGA
Ang4 FP	TGGTTGTGATTCCTCCAACTCT
Ang4 RP	TGTTGACATCTTTGCAAGGCG
fliC FP	CTAAACAAACTGGGTGGCGC
fliC RP	GCACCCAGGTCAGAACGTAA
bapA FP	ACGGTGACTACGCCTAAAAC
bapA RP	CTTCCGCCTCAGTCACTTTT
csgD FP	ACGATTATCCCTACCGTGAA
csgD RP	GCGGACTCGGTGCTGTTGTA
Salmonella 16S rRNA FP	GTGAGGTAACGGCTCACCAA
Salmonella 16S rRNA RP	TAACCGCAACACCTTCCTCC
sipC FP	GCAGTACGCCGCTACTCAGG
sipC RP	AGCGCGAATATTGCCTGCGA
pipB2 FP	TGGCGCCAATTTACAACACGC

pipB2 RP	CTCGCCTTTCGCAGGTTTGC

Table S3-List of Antibodies

Antibody Name	Catalogue Number
PV-1	A15906-20ul (Abclonal)
CD34	A19015-20ul (Abclonal)
Muc2	A4767 (Abclonal)
ZO-1	R26.4C DHSB
β actin	A3854 (Sigma)

GRAPHICAL ABSTRACT

O. splanchnicus bacteriocin

Flagella formation →

SPI-1 gene expression → Reduced *Salmonella* invasion

SPI-2 gene expression → Reduced *Salmonella* proliferation

GUT LUMEN

MUCUS

TJ/AJ

EPITHELIUM

TJ/AJ

GVB

LAMINA PROPRIA

Odoribacter splanchnicus colonization prior or post-*Salmonella* or *Listeria* infection

MUCUS

EPITHELIUM

Decreased TJ/AJ

Decreased TJ/AJ

PV-1

GVB

LAMINA PROPRIA

Salmonella or *Listeria* infection without *Odoribacter splanchnicus* colonization

Tight junctions/ Adherens junctions   PV-1 (Marker of Gut Vascular Barrier (GVB) disruption)

Odoribacter splanchnicus   *Salmonella* Typhimurium   *Listeria monocytogenes*   Non-flagellated *Listeria monocytogenes*

Odoribacter splanchnicus produced bacteriocin   *Salmonella* Typhimurium with reduced or no flagella