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Pro-angiogenic activity of TLRs and NLRs: a novel link between gut microbiota and intestinal angiogenesis

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Abstract

Background & Aims—In intestinal inflammation the gut microbiota induces an innate immune response by activating epithelial and immune cells that initiate or maintain inflammation. We investigated whether the microbiota can also activate local microvascular cells and induce angiogenesis.

Methods—Human intestinal microvascular endothelial cells (HIMEC) and intestinal fibroblasts (HIF) were exposed to bacterial ligands specific for TLR2/6 and 4, and NOD1 and NOD2, and cell proliferation, migration, transmigration, tube formation and production of pro-angiogenic factors were measured. The ability of the ligands to induce *ex vivo* vessel sprouting in an aortic ring assay and *in vivo* angiogenesis using a collagen gel assay were also assessed.

Results—Bacterial ligands induced proliferation, migration, transmigration, tube formation of HIMEC, vessel sprouting and *in vivo* angiogenesis; they also stimulated production of angiogenic factors from HIMEC and HIF, and HIF-derived angiogenic factors promoted HIMEC

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Author contribution <u>Anja Schirbel</u>: study concept and design; acquisition of data; drafting of the manuscript; critical revision of the manuscript for important intellectual content; obtained funding; analysis and interpretation of data; statistical analysis. <u>Sean Kessler</u>: study design; acquisition of data; critical revision of the manuscript for important intellectual content; drafting of the manuscript; technical support.

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proliferation. To various degrees, all ligands induced angiogenic responses, but these were ligandand cell type-dependent. Responses were mediated through RIP2-and TRAF6-dependent signaling, involved the MAPK and NF- κ B pathways and the upregulation of VEGF-R2 and FAK. Knockdown of RIP2 and TRAF6 by RNA interference and neutralization of IL-8, bFGF and VEGF inhibited TLR/NLR-induced HIMEC angiogenesis.

Conclusions—The gut microbiota can selectively activate mucosal endothelial and mesenchymal cells to promote specific angiogenic responses in a TLR- and NLR-dependent fashion. This innate immunity-mediated response may expand the mucosal microvascular network, foster immune cell recruitment, and contribute to chronic intestinal inflammation.

Keywords

angiogenesis; IBD; TLR; NLR

INTRODUCTION

Health depends on a mutually beneficial interaction between the host and its microbiota. It is increasingly evident that the commensal gut microbiota is involved not only in the development of systemic immunity¹, but also the pathogenesis of various immune-mediated conditions like type 1 diabetes, cardiovascular, neurologic and liver diseases^{2–5}. This is also true in inflammatory bowel disease (IBD), where microbial products play a role in initiation or maintenance of inflammation by modulating the function of epithelial and immune cells⁶. Non-immune cells, however, also contribute to IBD pathogenesis, such as endothelial and mesenchymal cells⁷, but whether and how microbial factors modulate their function is poorly understood.

Microbes are recognized by Toll-like (TLR) and NOD-like (NLR) receptors widely expressed by multiple cell types, leading to physiologic or pathological responses. In the intestine commensal bacteria can trigger inflammation in a host with a disrupted epithelial barrier^{8,9}, resulting in translocation and activation of cells in the mucosa. While the consequences of TLR and NLR activation on epithelial and immune cells have been extensively investigated^{10,11}, little information is available on the effect of microbial products on non-immune cells, particularly mucosal endothelial cells.

Microbial products may contribute to disease pathogenesis by affecting endothelial cell function in conditions such as atherosclerosis and liver diseases^{12,13}. This effect is mediated by specific ligands for TLRs or NLRs on endothelial cells^{14,15}, and subsequent triggering of angiogenesis, as reported in rheumatoid arthritis¹⁶. During postnatal development, intestinal angiogenesis accompanies the acquisition of gut microbiota¹⁷, but there is no information on whether activation of mature mucosal endothelial cells through TLRs/NLRs promotes angiogenesis, a component of human and experimental IBD^{18,19}.

Angiogenesis is a complex process involving not only endothelial cells but also mesenchymal cells, which are in close proximity of the microvasculature²⁰. Fibroblasts release various pro-angiogenic factors upon cytokine stimulation¹⁸ and express functional TLRs and NLRs²¹, but whether activation of these microbial receptors also induces release of such factors is unknown.

We investigated whether bacterial ligands induce angiogenesis by activating intestinal mucosa microvascular and mesenchymal cells. The results show that specific ligand-mediated activation of TLR2/6, TLR4, NOD1 and NOD2, which recognize a broad range of Gram-positive and -negative bacteria, promotes an angiogenic response mediated by a direct

stimulatory effect on endothelial cells and an indirect effect on mesenchymal cells leading to production of pro-angiogenic factors.

MATERIALS AND METHODS

Transfections

HIMEC were transfected with RIP2 or TRAF6 siRNA and their respective scrambled siRNA (both from Thermo Scientific, Lafayette, CO) using the Amaxa system (Amaxa Biosystems, Köln, Germany). HIMEC (0.5×10⁶) were suspended in 100 µl nucleofector solution for endothelial cells (Lonza, Walkersville, MD), mixed with different volumes of 20 nmol RIP2, TRAF6 or scrambled siRNA, transferred to a cuvette and nucleofected according the manufacturer's instructions. Cells were then transferred into 6-well plates containing pre-warmed culture medium. Some cells were left in nucleofector solution without applying electroporation to serve as negative control; 48 h after electroporation cells were used for the transmigration and MatrigelTM assays. Optimal doses and time points for optimal RIP2 and TRAF6 knockdown were pre-determined by immunoblot using antibodies against RIP2 (Alexis, San Diego, CA) and TRAF6 (Santa Cruz, Santa Cruz, CA), respectively (see supplemental materials).

HEK293T cells were transfected using Polyfect (Qiagen, Valencia, CA) according to the manufacturer's instructions. Briefly, a transfection mix comprised of 0.25 µg pxB-EGFP (gift of Xin Lin, University of Texas)²²,15 ng pcDNA3-HA-Nod1 (gift of Gabriel Nuñez, University of Michigan²³, 1.75 µg pcDNA3 and 12 µL Polyfect was added to 2×10^5 HEK293T cells in a 6 well plate. Medium was changed to fresh DMEM with 10% fetal bovine serum (FBS) after 5h and cells used for *in vitro* MatrigelTM assays 48 h post-transfection.

Mouse aortic ring assay

Rings of mouse aorta were cultured in three-dimensional collagen gels as described²⁴. Tie2green fluorescent protein (GFP) mice (Jackson Laboratories, Bar Harbor, ME) expressing GFP exclusively in endothelial cells were sacrificed by CO₂ inhalation, and aortas dissected and transferred to ice-cold minimum essential medium (MEM, Sigma-Aldrich, St. Louis, MO) containing 1% pen/strep/fungizone (PSF, Lonza). The peri-aortic fibroadipose tissue was carefully removed and 1 mm long aortic rings were sectioned and rinsed in MEM. Ring-shaped explants of mouse aorta were then embedded in 300 μl of a mixture of growth factor-depleted MatrigelTM (BD Biosciences, Bedford, MA) and 199 medium containing 1% PSF and 2.5% mouse serum (Innovative Research, Novi, MI) in 48-well plates. After 20 min polymerization at 37°C 500 μl 199 medium supplemented with 2.5% mouse serum, 1% PSF, with and without bacterial ligands or VEGF, were added. The cultures were kept at 37°C for two weeks, medium was changed every second day, and gels were examined every other day by phase microscopy and harvested at predetermined time points.

Harvested aortic rings in MatrigelTM were incubated overnight in Histochoice (Amresco, Solon, OH) prior to embedding in paraffin. Slide sections were deparaffinized and rehydrated by processing through Clear-rite 3 (2×3 minutes), Flex-100 (2 min, then 1 min) and Flex-95 (2 min, then 1 min) (Richard-Allan Scientific, Kalamazoo, MI) followed by tap water rinse. Tissue sections were blocked in HBSS with 2% FBS for 1 hr at 25 C°. Primary antibodies, (rabbit anti-CD31 and goat anti-GFP 1:100 dilution; Abcam 28365 and 6673, respectively; Abcam, Cambridage, MA) were applied in the same blocking buffer overnight at 4 C°. Slides were washed twice in HBSS and once in blocking buffer prior to applying donkey anti-rabbit Alexa-Fluor 488 (Life Technologies, Grand Island, NY) diluted 1:1000 (Alexa A-21206). Slides were washed 3 times prior to the addition of rabbit anti-goat Biotin

(Jackson Immunological Res 305-066-045) diluted1:100 and Streptavidin-568 (Alexa S-11226) diluted 1:500 in blocking buffer. Slides were mounted in Vectashield plus DAPI (Vector Labs, Burlingame, CA) and examined with a fluorescence microscope at $40 \times$ magnification.

In vivo collagen gel assay

Collagen gels were prepared as described²⁵. A solution comprising 1.5 mg/ml rat tail collagen Type 1 (BD Biosciences), 25 mM HEPES (Lonza), 1.5 mg/ml sodium bicarbonate, 10% FBS, 30% EGM-2 medium (Lonza) in EBM-2 medium, at pH 7.4 was prepared on ice. After gentle mixing 1 ml of the suspension was placed into 12-well plates, allowed to polymerize for 30 min at 37°C, and covered with complete EGM-2 overnight. Gels were bisected and implanted into a preformed subcutaneous pocket lateral of the midventral line of mice.

Gels contained medium alone, VEGF (200 ng/ml; R&D Systems, Minneapolis, MN) or one of the following TLR4 or NOD1 ligands: ultrapure lipopolysaccharide (LPS) from 0111:B4 *E.coli* (100 ng/ml; InvivoGen, San Diego CA), crude LPS from 0111:B4 *E.coli* (1 μ g/ml; Sigma L2630) or iE-DAP (5 μ g/ml; AnaSpec, Inc., San Jose, CA). Seven days after gel implantation mice were sacrificed by CO₂ inhalation and gels were excised, fixed in 10% formaldehyde for 4 hours followed by 70% ethanol overnight, paraffin embedded, stained for H&E and anti-mouse CD31 (Abcam).

RESULTS

NLR and TLR activation induce HIMEC migration, transmigration and proliferation

Angiogenesis involves a series of cellular events among which growth and movement are essential to new vessel formation²⁶. We tested whether endothelial cell migration, transmigration and proliferation were induced by TLR or NLR activation. Using the wound migration assay, ligands for TLR2/6, TLR4, NOD1 and NOD2 significantly (p<0.04-0.001) enhanced migration of HIMEC regardless of whether control or IBD HIMEC were used (Fig.1A and **supplemental Fig. 1A and 1B**). Exposure to basic fibroblast growth factor (bFGF), but not TNF- α , both potent non-microbial stimuli, significantly increased migration (Fig.1A). In contrast, transmigration was significantly increased only by TLR4 and NOD1 activation (p<0.03-0.01) (Fig. 1B), and only IBD HIMEC significantly increased proliferation in response to NOD1 activation (p<0.01) (Fig. 1C). As seen for migration, TNF- α failed to significantly enhance transmigration and proliferation. IBD cells tended to migrate more than control, but this difference was not significant; the same was true in regard to transmigration and proliferation.

Induction of HIMEC tube formation and vessel sprouting by NLR and TLR activation

To determine whether TLRs and NLRs could mediate angiogenesis under conditions representative of what may occur in tissue, we performed assays involving tube formation and vessel sprouting *ex vivo* and *in vivo*.

Initially we tested the ability of TLRs and NLRs to mediate tube formation in MatrigelTM, a more physiological response than that resulting from monolayer culture²⁷. Compared to medium alone, HIMEC stimulated with TLR2/6, TLR4, NOD1 or NOD2 ligands readily formed tubules with thinner and longer bridges; this effect was comparable to or faster than that induced by VEGF, while TNF-a failed to induce tube formation (Fig. 2A and **supplemental Fig. 2**).

Second, we performed a murine aortic ring assay, which combines the advantages of both *in vivo* and *in vitro* assays and is based on induction of endothelial cell branching²⁸. Similarly to tube formation, all ligands stimulated vessel sprouting (Fig. 2B), activation of TLR4 and NOD1 being consistently more effective than TLR2/6 and NOD2 activation, and as effective as VEGF. The endothelial nature of the sprouting cells was indicated by typical microvascular structures with multiple subsequent branching (**Supplemental Fig. 3A**), and confirmed by CD31 staining of aortic rings from Tie2 GFP mice (**Supplemental Fig. 3B and 3C**). No differences were noted between control and IBD HIMEC in regard to tube formation or cell sprouting.

Next, the capacity of TLR4 and NOD1 ligands to induce angiogenesis *in vivo* was tested using a collagen gel system²⁵. Seven days after implantation, control gels were essentially devoid of cells or vessels, and VEGF-containing gels only contained scattered cells. In contrast, gels containing ultrapure LPS or NOD1L exhibited various degrees of cell infiltration and microvascular formation compared to control gels; when crude LPS, which contains various other bacterial products²⁹, was used gels displayed heavy cellular infiltration and the highest degree of *in vivo* angiogenesis (Fig. 3).

Induction of HIMEC proliferation by TLR/NLR-activated HIF conditioned medium

Angiogenesis involves the interaction of endothelial cells with other cell types²⁶. Since mesenchymal cells are in close proximity of endothelial cells and produce pro-angiogenic factors^{30–32}, we investigated whether mucosal fibroblasts contribute to mucosal angiogenesis. HIF were exposed to the same TLR and NLR ligands and the ability of HIF-conditioned medium to induce HIMEC angiogenesis was tested.

We first cultured HIF alone, in the presence of TNF- α , or individual TLR2/6, TLR4, NOD1 and NOD2 ligands; after 24 hours medium was removed, cells washed to remove any residual stimuli, and cultured alone for an additional 48 h to generate HIF-conditioned medium. Conditioned medium from both unstimulated and TNF- α -stimulated HIF dosedependently increased HIMEC proliferation with maximal effect at 50% dilution (**Supplemental Fig. 4A**). At this dilution unstimulated and ligand-activated HIF-conditioned medium induced a significant (p<0.05–0.03) increase in HIMEC proliferation compared to the direct effect of the medium or ligands alone (Fig. 4A). HIF medium conditioned by bFGF or IL-1 β induced a similar response (not shown). This effect was not induced by FBS, as HIMEC exposed to various concentrations of FBS for variable periods of time failed to proliferate more than HIMEC cultured in the absence of FBS (legend of Fig. 4A). In view of the potent stimulatory activity of HIF-conditioned medium even in the absence of added stimuli, only TLR4 activation induced a significant (p<0.03) further increase of HIMEC proliferation compared to medium of HIF cultured alone (Fig. 4A). Results were independent of the control or IBD origin of HIMEC or HIF.

Induction of HIMEC and HIF pro-angiogenic factors by TLR/NLR activation

HIMEC produce autocrine IL-8 to promote angiogenesis¹⁸, but induction of HIMEC proliferation by HIF-conditioned medium suggested the presence of pro-angiogenic factors in this medium^{30–32}. Therefore, we investigated the production of IL-8, bFGF and VEGF by HIMEC and HIF in response to TLR/NLR activation as well as TNF- α , bFGF and VEGF. Bacterial ligands stimulated production of angiogenic factors, but this response was ligand-or cell type-dependent.

HIMEC production of IL-8 increased significantly in response to TLR4, NOD1 and NOD2 but not TLR2/6 ligands (Fig. 4B); bFGF increased significantly only in IBD HIMEC responding to NOD1 and NOD2 ligands (**Supplemental Fig. 4B**), while VEGF was not

modulated (not shown). HIF produced significantly more IL-8 in response to TLR2/6, TLR4 and NOD1 but not NOD2 ligands (Fig. 4C); HIF spontaneously produced VEGF, but its level did not increase in response to TLR/NLR activation (not shown). When compared to bacterial ligands, TNF-a induced HIMEC to produce IL-8 to a degree comparable to that induced by TLR4 activation, but higher than NOD1 and NOD2 activation (Fig. 4B), and TNF-a was far more potent than TLR/NLR ligands in stimulating IL-8 production by HIF (Fig. 4C). bFGF and VEGF exhibited no or low capacity to induce angiogenic factor production by HIMEC or HIF. Significant differences were not noted between control and IBD HIMEC.

In additional experiments the following angiogenic factors were assessed in HIF supernatants: insulin-like growth (IGF-1), IL-6, placenta growth factor-1 (PIGF-1), stem cell factor (SCF), TGF- β (transforming growth factor-1), epidermal growth factor (EGF), nerve growth factor (NGF), leptin, PDGF-BB (platelet-derived growth factor-BB), monocyte chemoattractant protein-1 (MCP-1), and macrophage inflammatory protein-1a (MIP-1a); of these, variable amounts (from 16 to 2500 pg/ml) of IGF-1, IL-6, PDGF-BB, MCP-1 and MIP-1a were present in the supernatants; PIGF-1, SCF, TGF- β , EGF, NGF and leptin were either absent or present at borderline detection levels.

Inhibition of angiogenesis by blockade of TLR/NLR-induced angiogenic factors

We next investigated to what extent the TLR/NLR-induced angiogenic response was a direct effect of the bacterial ligands or an indirect effect dependent on production of angiogenic factors. IL-8 was blocked together with bFGF or VEGF and HIMEC migration and proliferation were measured as indicators of angiogenesis (Fig. 1A, 1C).

The combined blockade of bFGF and IL-8, both of which are produced by HIMEC (see above), significantly (p<0.05–0.001) reduced HIMEC migration directly induced by TLR/ NLR activation (**Supplemental Fig. 5A**); bFGF-, but not TNF-a-, dependent migration was also reduced. In these experiments proliferation was unaffected (not shown).

We then cultured HIMEC in HIF-conditioned medium with and without blockade of IL-8 and VEGF and measured cell proliferation. A significant (p<0.05-0.01) reduction of proliferation was observed in response to TLR2/6-, TLR4- and NOD1- but not NOD2- conditioned medium (supplemental results and **Supplemental Fig. 5B**). Since TNF-a induced abundant IL-8 production by HIF (Fig. 4C), neutralization also significantly decreased the proliferation induced by TNF-a-conditioned medium; the effect of control bFGF was not significantly decreased. Results were independent of the control or IBD origin of the HIMEC.

Dependency on RIP2 for NLR-induced angiogenesis

RIP2 is an adaptor molecule essential for NLR signaling³³, and we investigated its involvement in NOD1- and NOD2-mediated angiogenesis by RNA interference. After determining optimal knockdown conditions (**Supplemental Fig. 6A**), 0.1 nmol/ml and 48 h were selected and applied to transmigration and tube formation assays. Compared to scrambled siRNA, RIP2 siRNA significantly (p<0.01) reduced transmigration of HIMEC stimulated by NOD1 and NOD2 ligands, but not by bFGF or TNF- α (Fig. 5A). Comparable results were observed in the tube formation assay, with marked inhibition seen when NOD1 or NOD2 ligands were used (Fig. 5B). A strong inhibition of tube formation was unexpectedly observed in HIMEC transfected with RIP2 siRNA even in the absence of NLR ligands (Fig. 5B). Therefore, we investigated whether NOD1 or NOD2 ligands were inadvertently present in the MatrigelTM itself, but this was not the case as shown by negative fluorescence-based NF- κ B reporter gene assays (supplemental Results and Supplemental Fig. 7).

Involvement of TRAF6 in TLR/NLR-induced angiogenesis

Since TRAF6 is an adaptor molecule essential for both NLR and TLR signaling³⁴, we blocked its activity with TRAF6 siRNA to investigate its involvement in TLR2/6-, TLR4-, NOD1- and NOD2-mediated angiogenesis. After establishing optimal inhibitory conditions (0.4 nmol/ml and 48 h) (supplemental Results and **Supplemental Fig. 6B**), transmigration and tube formation assays were performed. Knockdown of TRAF6 mRNA significantly (p<0.04–0.006) inhibited all TLR/NLR-induced HIMEC transmigration compared to cells transfected with scrambled siRNA, with no significant effect on bFGF-induced transmigration (Fig. 6A). Using TRAF6 siRNA tube formation was inhibited in TLR- and NLR-activated HIMEC, but not in HIMEC cultured in medium alone (Fig. 6B).

Involvement of the MAPK and NF-κB pathways, VEGF-R2 and FAK in TLR/NLR-induced angiogenesis

Multiple pathways are involved in signaling downstream of TLR and NLR^{35,36}, and we investigated the activation of MAPK and NF- κ B pathways in HIMEC by assessing the phosphorylation of IKBa, p38 and ERK. As shown in Fig. 7A, activation of TLR2/6 and TLR4 as well as of NOD1 and NOD2 by their respective ligands led to a strong and time-dependent phosphorylation of all the above signaling molecules.

We finally investigated additional molecules critically involved in events required for angiogenesis: these included VEGF-R2, the main receptor mediating the angiogenic activity of VEGF, and FAK, which is broadly involved in cell spreading and migration^{37,38}. Compared to unstimulated cells, all ligands upregulated the expression of VEGF-R2 to a degree comparable to that induced by TNF-a, and induced its phosphorylation to an even higher degree (Fig. 7B); the same ligands induced phosphorylation of FAK similar to or greater than that induced by VEGF, while levels of total FAK remained unchanged (Fig. 7C). In all experiments investigating RIP2, TRAF6, MAPK and NF-kB, VEGF-R2 and FAK no differences were observed between control and IBD HIMEC.

DISCUSSION

This study demonstrates that microbial products can directly activate mucosal endothelial cells and induce proliferation, migration, transmigration, tube formation and vessel sprouting. These responses are mediated through various TLRs and NLRs and promote angiogenesis. Additionally, microbial products contribute indirectly to angiogenesis by inducing the TLR- and NLR-dependent production of pro-angiogenic factors by mucosal mesenchymal cells. Together, these observations suggest that the intestinal microbiota contributes to angiogenesis during intestinal inflammation^{18,19}.

The type and degree of TLR2/6-, TLR4-, NOD1- and NOD2-mediated responses depended primarily on the effect induced without significant differences between control and IBD HIMEC. Activation of TLR4 and NOD1 was particularly effective in inducing transmigration, tube formation and *ex vivo* and *in vivo* vessel sprouting, while TLR2/6 and NOD2 activation had the strongest effect on migration. NOD2 ligation was relatively less effective, probably due to lower expression or affinity in HIMEC, as suggested by lower mRNA expression compared to NOD1 mRNA (unpublished). Overall responses were comparable to those induced by bFGF or VEGF, two of the most powerful pro-angiogenic factors. Therefore, it appears that TLRs and NLRs better promote endothelial cell movement than growth. Additional to these *in vitro* observations, the collagen gel assays confirmed that

single ligands specific for TLR4 and NOD1 are sufficient to induce angiogenesis *in vivo*. Crude LPS, which contains other bacterial products²⁹, was more potent that ultrapure LPS, suggesting additive or synergistic effects in angiogenesis induction, a response likely to reflect the actual *in vivo* situation.

We also tested the pro-angiogenic activity of TNF- α , a prototypical inflammatory cytokine abundantly produced in IBD³⁹. The role of TNF- α as a pro-angiogenic mediator is controversial because few studies claim that TNF- α exerts pro-angiogenic activity *in vivo*, whereas the majority of reports fail to find such activity *in vito*⁴⁰. TNF- α or TNF- α conditioned medium failed to induce HIMEC migration, transmigration or proliferation, and did not promote tube formation. Conversely, TNF- α was at least as effective as TLR and NLR ligands in inducing IL-8 production by HIMEC and HIF and upregulating VEGF-R2 expression on HIMEC. This suggests that microbial stimuli may be more effective than inflammatory cytokines in the direct induction of mucosal angiogenesis; inflammatory cytokines, like TNF- α , may complement this effect indirectly by priming endothelial cells, stimulating mesenchymal cells to secrete angiogenic factors, and upregulating their receptor on the nearby microvasculature.

TLR and NLR activation robustly stimulated HIMEC movement, and this was mediated by a time-dependent phosphorylation of signaling molecules of the MAPK and NF-κB pathways, including p38, ERK and IKBa. In addition, TLR and NLR activation led to phosphorylation of FAK, a tyrosine kinase essential to integrin-mediated cell adhesion⁴¹. FAK phosphorylation is indispensable to cell spreading, migration, survival and cycle progression and it has been recently shown to also promote angiogenesis³⁸. TLR and NLR activation also upregulated VEGF-R2, the main receptor through which VEGF exerts proangiogenic activity³⁷. Together, the activation of the MAPK and NF-kB pathways, the induction of p-FAK, and of VEG-R2 mimic the action of classical pro-angiogenic factors, supporting the notion that microbial products are important mediators of intestinal angiogenesis acting through selective TLRs and NLRs.

Next we investigated the role molecules traditionally involved in TLR and NLR signaling, including the adaptor RIP2 for NLRs and TRAF6 for TLRs and NLRs^{33,34}. RIP2 knockdown significantly inhibited NOD1- and NOD2-induced transmigration and tube formation. Unexpectedly, RIP2 knockdown inhibited tube formation of HIMEC seeded in medium alone, a response not due to the inadvertent presence of NOD ligands in medium or MatrigelTM. Thus, RIP2 exerts functions not exclusively related to NLR signaling, as indicated by reports showing that RIP2 may be involved in cell death and survival in Schwann cells, T cell receptor-mediated NF-κB activation, and myogenic responses^{42–44}. The involvement of TRAF6 was also evidenced when its knockdown inhibited transmigration and tube formation. So, TLR/NLR signaling pathways acting in immune and epithelial cells are also functional in mucosal microvascular endothelial cells with a cell-type specific outcome, i.e., promotion of intestinal angiogenesis.

Leaking of microbial products in the inflamed intestine allows interaction with mucosal cells bearing their specific receptors⁴⁵, including endothelial and mesenchymal cells^{14,15,21}. That endothelial cells, including HIMEC, produce their own pro-angiogenic factors acting in an autocrine fashion and gut mucosal extracts contain pro-angiogenic factors is known^{18,19}. However, the ability of mucosal mesenchymal cell-derived factors to act in a paracrine pro-angiogenic fashion was unknown. Medium conditioned by HIF alone was able to induce HIMEC proliferation, which increased further when HIF were pre-stimulated with LPS. Indeed, HIF produced vast amounts of IL-8 in response to TLR2/6 activation, and increased amounts in response to TLR4 and NOD1 activation. HIF also produced IGF-1, IL-6, PDGF-BB, MCP-1 and MIP-1a, all of which likely act in a complementary fashion to promote

mucosal angiogenesis. Blocking some of these pro-angiogenic factors in HIF-conditioned medium inhibited TLR/NLR-induced HIMEC proliferation and migration, demonstrating a crucial role of fibroblasts in inflammation-induced angiogenesis comparable to that seen in neoplasia-induced angiogenesis³¹.

In summary, this study shows that different microbiota-derived products induce mucosal endothelial and mesenchymal cells to promote intestinal angiogenesis through the selective induction of specific pro-angiogenic pathways. Angiogenesis fosters recruitment of immune cells and sustains inflammation, and therapeutic manipulation of the gut microbiome may prevent or attenuate chronic intestinal inflammation.

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Abbreviations

bFGF	basic fibroblast growth factor
FAK	focal adhesion kinase
HIF	human intestinal fibroblast
HIMEC	human intestinal microvascular endothelial cell
NLR	Nod-like receptor
TLR	Toll-like receptor
TRAF6	TNF receptor-associated factor 6
RIP2	receptor interacting protein-2
VEGF	vascular endothelial growth factor
VEGF-R2	vascular endothelial growth factor receptor 2

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Figure 1.

A) TLR and NLR activation-induced HIMEC migration

Wounded control and IBD HIMEC monolayers were exposed to optimal stimulatory doses of bFGF, TNF- α and TLR2/6L, TLR4L, NOD1 and NOD2 ligands, and migrated cells counted after 24 h. *p<0.04–0.01 for stimulated *vs* unstimulated (medium) control HIMEC; **p<0.02–0.002 for stimulated *vs* unstimulated (medium) IBD HIMEC (n=10 and 14 for control and IBD HIMEC, respectively); L: ligand; hpf: high power field.

B) TLR and NLR activation-induced HIMEC transmigration

Control and IBD HIMEC were place in the upper compartment of a Boyden chamber, optimal stimulatory doses of bFGF, TNF-a and TLR2/6L, TLR4L, NOD1 and NOD2 ligands were placed in the lower compartment, and the number of transmigrated cells counted after 8 h. *p<0.03–0.01 for stimulated *vs* unstimulated (medium) control HIMEC; **p<0.01 for stimulated *vs* unstimulated (medium) IBD HIMEC (n=8 for both control and IBD HIMEC).

C) TLR and NLR activation-induced HIMEC proliferation

Control and IBD HIMEC monolayers were exposed to optimal stimulatory doses of bFGF, TNF- α and TLR2/6L, TLR4L, NOD1 and NOD2 ligands, and proliferation assessed by ³H thymidine uptake after 24 h. *p<0.001 for stimulated *vs* unstimulated (medium) control HIMEC; **p<0.01–0.001 for stimulated *vs* unstimulated IBD HIMEC (n=10 and 11 for control and IBD HIMEC, respectively).



Figure 2.

A) TLR and NLR activation-induced HIMEC tube formation

Control and IBD HIMEC were seeded on Matrigel[™] in the presence and absence of optimal stimulatory doses of VEGF, TNF-a and TLR2/6L, TLR4L, NOD1 and NOD2 ligands, and tube formation visually assessed up to 24 h. Figure is representative of 9 and 8 experiments for control and IBD HIMEC, respectively, at 6 h.

B) TLR- and NLR-induced HIMEC vessel sprouting in murine aortic ring assay Murine aortic rings were placed in MatrigelTM in the presence and absence of optimal stimulatory doses of VEGF, TNF- α and TLR2/6L, TLR4L, NOD1 and NOD2 ligands, and vessel sprouting visually assessed after 6 days. Figure is representative of 5 and 5 experiments for control and IBD HIMEC, respectively.



Figure 3.

VEGF-, TLR4L- and NLRL-induced cell infiltration and vessel formation *in vivo* Collagen gels individually containing medium alone (control), VEGF, TLR4L, or NOD1L were surgically implanted in the abdominal subcutaneous tissue of mice, harvested after 7 days, and stained for H&E and CD31 to visualize vessel growth. Upper panels were taken at 10× and lower panels at 40× magnification. Figure is representative of 5 experiments (TLR4L-UP: ultrapure LPS; TLR4L-C: crude LPS).



Figure 4.

A) HIMEC proliferation in response to HIF-conditioned medium

HIMEC were cultured alone (medium), in the presence of TNF- α or optimal stimulatory doses of TLR and NLR ligands (black columns), or in HIF medium conditioned alone (medium), by TNF- α , or the same doses of TLR and NLR ligands (grey columns); after 24 h proliferation was assessed by ³H thymidine uptake. *p<0.05–0.03 for TLR/NLR activation-conditioned HIF medium compared to direct stimulation with TLR and NLR ligands; **p<0.03 for TLR4 activation-conditioned HIF medium compared to unstimulated HIF medium (n=10 for both control and IBD HIMEC). HIF (n=3) were also cultured in the presence of 0%, 2%, 5%, 10% and 20% FBS for 24, 48 and 72 hs, respectively, and proliferation assessed as described above. None of the different concentrations of FBS induced a significant increase in HIMEC proliferation compared to HIMEC cultured alone (0% FBS) at any of the 3 times points (not shown).

B) IL-8 production by TLR- and NLR-activated HIMEC

Control and IBD HIMEC monolayers were exposed to optimal stimulatory doses of VEGF, TNF-a, TLR2/6, TLR4, NOD1 and NOD2 ligands for 24 h and levels of IL-8 measured by ELISA in the culture supernatants. *p<0.03-0.001 for stimulated *vs* unstimulated (medium) control HIMEC; **p<0.004-0.001 for stimulated *vs* unstimulated (medium) IBD HIMEC (n=5 for both control and IBD HIMEC).

C) IL-8 production by TLR- and NLR-activated HIF

Control and IBD HIF monolayers were exposed to optimal stimulatory doses of bFGF, TNF-a, TLR2/6, TLR4, NOD1 and NOD2 ligands for 24 h and levels of IL-8 measured by ELISA in the culture supernatants. *p<0.05–0.004 for stimulated *vs* unstimulated (medium) control HIF; **p<0.03–0.004 for stimulated *vs* unstimulated IBD HIF (n=3 and 4 for control and IBD HIF, respectively).



Figure 5.

A) Effect of RIP2 knockdown on NLR activation-induced HIMEC transmigration
Transmigration was performed as described above in the legend of Figure 1B, using cells
transfected with scrambled or RIP2 siRNA and then exposed to medium alone, bFGF, TNF-a, or NOD1 and NOD2 ligands for 8 h. *p<0.01 for siRIP2 RNA vs scrambled siRNA (n=3 and 3 for control and IBD HIMEC, respectively); hpf: high power field.
B) Effect of RIP2 knockdown on NLR-induced HIMEC tube formation
Tube formation assay was performed as described above in the legend of Figure 2A, using cells transfected with scrambled or RIP2 siRNA and then exposed to medium alone, NOD1L or NOD2L. Figure is representative of 3 and 2 experiments with control and IBD HIMEC, respectively, at 4 h.



Figure 6.

A) Effect of TRAF6 knockdown on TLR and NLR activation-induced HIMEC transmigration

Transmigration was performed as described above in the legend of Figure 1B, using cells transfected with scrambled or TRAF6 siRNA and then exposed to medium alone, bFGF or TLR2/6, TLR4, NOD1 and NOD2 ligands for 8 h. *p<0.04–0.006 for siTRAF6 RNA *vs* scrambled RNA. (n=3 and 3 for control and IBD HIMEC, respectively); hpf: high power field.

B) Effect of TRAF6 knockdown on TLR- and NLR-induced HIMEC tube formation Tube formation assay was performed as described above in the legend of Figure 2A, using cells transfected with scrambled or TRAF6 siRNA and then exposed to medium alone, TLR2/6, TLRL4, Nod1L or Nod2L. Figure is representative of 3 and 2 experiments with control and IBD HIMEC, respectively, at 5 h.





Figure 7.

A) TLR- and NLR-mediated MAPK and NF-rB activation

HIMEC monolayers were exposed to optimal stimulatory doses of TLR2/6, TLR4, NOD1 or NOD2 ligands for the indicated time points, harvested, lysed, and protein extracted for immunoblotting using antibodies against total and phosphorylated (p) p38, ERK, IKBa and GAPDH. Figure is representative of 3 control and 3 IBD HIMEC, respectively. B) TLR and NLR activation-induced VEGF-R2 phosphorylation

HIMEC monolayers were exposed to optimal stimulatory doses of TNF-a or TLR2/6, TLR4, NOD1 or NOD2 ligands for 48 h, harvested, lysed, and protein extracted for

immunoblotting using the respective antibody and GAPDH antibody. Lower panel shows densitometric analysis of the ratio of p-VEGF-R2/VEGF-R2 under each condition. Figure is representative of 3 control and 5 IBD HIMEC.

C) TLR and NLR activation-induced FAK phosphorylation

HIMEC monolayers were exposed to optimal stimulatory doses of VEGF or TLR2/6, TLR4, NOD1 or NOD2 ligands for 48 h, harvested, lysed, and protein extracted for immunoblotting using the respective antibody and GAPDH antibody. Lower panel shows densitometric analysis of the ratio of p-FAK/total FAK under each condition. Figure is representative of 3 control and 5 IBD HIMEC.