Evidence is accumulating that chronic inflammation contributes to carcinogenesis, including gastrointestinal cancers, lung cancer, and prostate cancer. Chronic colitis-associated colorectal carcinogenesis (CAC) is a typical model to study inflammation-associated carcinogenesis. Ulcerative colitis (UC), a major form of inflammatory bowel disease (IBD), is characterized by a long-lasting cycle of remission and exacerbations of ulceration of bowels, abdominal pain, diarrhea, bloody stool, and other systemic symptoms. It is reported that patients suffering from UC for more than 10 years are 6 to 10 times more susceptible to develop colorectal cancer than the general population. Furthermore, both the severity and the duration of the chronic IBD significantly correlate with the ...
Risk of colorectal cancer. Although inflammatory cytokines that mediated signaling in linking IBD and CAC were recently reported, such as tumor necrosis factor-\(\alpha\) (TNF-\(\alpha\))/inhibitor of nuclear factor kappa-B kinase-\(\beta\) (IKK-\(\beta\))/NF-\(\kappa\)-B and vascular endothelial growth factor (VEGF)/VEGF receptor (VEGFR), the underlying mechanisms need to be further explored.

The occurrence and progression of CAC are believed to be an orchestrated process involving the complex interplay between premalignant epithelial cells and the inflammatory environment, where blood vessel formation and inflammatory immune cell infiltration may play fundamental roles. Therefore, targeting angiogenesis and lymphocyte infiltration become promising therapeutic strategy to alleviate inflammation. Ideally, a simultaneous blockade of lymphocyte infiltration and angiogenesis by targeting one molecule would be more efficient for modulating inflammation and preventing CAC.

CD146, an adhesion molecule belonging to the immunoglobulin superfamily, has been reported to be expressed on certain cancer cells, endothelial cells, and some subpopulations of leukocyte cells. Our previous studies have identified that CD146 plays a key role in tumor-associated angiogenesis, as well as in tumor metastasis. The CD146 antibody AA98 inhibits tumor angiogenesis, respectively, reported increased expression of CD146 in patients with IBD. However, the function of CD146 and its regulatory mechanisms in IBD remain unknown. There are also clinical observations reporting that CD146 expression was associated with other inflammatory diseases such as rheumatoid arthritis, chronic renal failure, and diabetes through mechanisms yet unclear.

Therefore, it is of great interest to study how CD146 functions in inflammatory diseases, and more importantly, its role in chronic inflammation-associated carcinogenesis.

In this study, we found that endothelial CD146 plays important roles in the progression of IBD and CAC. CD146 was overexpressed on the endothelial cells in colitis and CAC model mice, but not on epithelial cells and tumor cells. Eliminating endothelial CD146 by conditional knockout or blocking CD146 by antibody AA98 significantly alleviated the disease severity in murine models of colitis and decreased tumor incidence in a murine model of CAC. Mechanistic study revealed that TNF-\(\alpha\) and IL1-\(\beta\) strongly induced CD146 expression on endothelial cells by direct NF-\(\kappa\)-B transactivation. Up-regulated endothelial CD146 played a dual function in inflammation, that is, promoting angiogenesis and recruiting lymphocytes, especially proinflammatory Th1 and Th17 cells. This, in turn, results in a vicious cycle of chronic inflammation, which promotes the occurrence of CAC. Our study suggests that CD146 is an important molecule involved in inflammation and carcinogenesis, which can be a therapeutic target in the treatment of IBD and prevention of CAC.

Materials and Methods

Antibodies and Reagents

The following antibodies were used in this study: anti-CD146 monoclonal antibody AA98, anti-mTNF-\(\alpha\) monoclonal antibody V1q, used as functional blocking antibody in animal experiments. Anti-p-Akt, anti-p-p65, anti-p-MAPKAPK-2, anti-p65 (Cell Signaling Technology, Danvers, MA), anti-glyceraldehyde-3-phosphate dehydrogenase (GAPDH) (Santa Cruz Biotechnology, Santa Cruz, CA), antibody AA98, and horseradish peroxidase-conjugated anti-rabbit secondary antibodies (GE Healthcare, Chalfont St Giles, UK) were used in the Western blot assays. Anti-human and anti-mouse CD31 (Cymbus Biotechnology, Chalmers Ford, UK), anti-mouse CD146 (clone 9F1), anti-mouse CD45, anti-mouse CD4 (BioLegend, San Diego, CA), anti-\(\beta\)-catenin (BD Biosciences, San Jose, CA), anti-mouse IL17 (clone 17F3; ebioScience, San Diego, CA), anti-Ki-67 (clone M-19; Santa Cruz Biotechnology), and DAPI (AppliChem, Darmstadt, Germany) were used in immunohistochemistry or immunofluorescence analysis. Anti-p65 polyclonal antibody and Isotype-matched rabbit IgG (Abcam) were used in the chromatin immunoprecipitation (ChIP) assay. Phycocerythin (PE)-conjugated AA98 (labeled by Tianjin SunGene Biotech, Tianjin, China), fluorescein isothiocyanate-conjugated anti-mouse IL17A (clone 17F3), and alphalocysocyanin (APC)-conjugated anti-mouse CD4 (clone GK1.5) (Tianjin SunGene Biotech), PerCP cy5.5-conjugated anti-mouse CD3, and PE-conjugated anti-mouse interferon-\(\gamma\) (IFN-\(\gamma\)) (ebioScience) were used in flow cytometry assays. Anti-CD146 siRNA and control green fluorescent protein (GFP) siRNA were synthesized by Invitrogen (Carlsbad, CA) using sequences as previously reported. P65 siRNA was purchased from Santa Cruz Biotechnology. The recombinant cytokines TNF-\(\alpha\), IFN-\(\gamma\), IL1-\(\beta\), and IL6 were purchased from PeproTech (Rocky Hill, NJ). Phytohaemagglutinin-P and lipopolysaccharide (LPS) were purchased from Sigma-Aldrich (St. Louis, MO). Dextran sulfate sodium (DSS) (mol. wt. 36,000 to 50,000) and azaoyxymethane (AOM) were obtained from MP Biomedicals (Santa Ana, CA) and Sigma-Aldrich, respectively. A BD Cytometric Bead Array kit for flow cytometry—based detection of serum cytokines was purchased from BD Biosciences. A TUNEL in situ apoptosis detection kit was purchased from TaKaRa Bio (Otsu, Japan). Human colon disease spectrum tissue microarray (BC05002) was bought from US Biomax (Rockville, MD).

Cell Culture and Animals

Human umbilical vein endothelial cells (HUVECs) were obtained from CellSystems Biotechnologie Vertrieb (St. Katharinen, Germany). Mouse endothelioma cell line used was sEnd1.1. Human monocyte cell-line THP-1 cells were
obtained from ATCC (Manassas, VA). These cells were cultured in RPMI 1640 medium containing 10% fetal calf serum at 37°C with 5% CO₂.

All animal experiments were performed in compliance with the Guidelines for the Care and Use of Laboratory Animals and were approved by the Biomedical Research Ethics Committee of the Institute of Biophysics, Chinese Academy of Sciences. C57BL/6J mice were obtained from the Animal Center of the Chinese Academy of Medical Science, Beijing. Tek⁺/CreCD146loxP/loxP mice were generated using a Cre/loxP recombination system. One loxP site was inserted in the intron before the promoter region of the CD146 (MCAM) gene, and one loxP site was inserted in the intron after the first exon. The Cre gene was driven by the promoter Tek of the Tie2 gene, which is expressed specifically on endothelial cells. Tek⁺/CD146loxP/loxP mice were backcrossed to mice on a C57BL/6J background for a minimum of nine generations. Tek⁺/Tek⁻/CD146loxP/loxP mice were then crossed with Tek⁺/CD146loxP/loxP mice. The F1 mice with genotypes Tek⁺/Tek⁻/CD146loxP/loxP were crossed with Tek⁺/+-CD146loxP/loxP mice to obtain Tek⁺/Tek⁻/CD146loxP/loxP [named wild-type (WT)] mice. The F1 mice to obtain CD146EC-KO and their WT counterpart, or C57BL/6J mice were injected i.p. with 12.5 mg/kg AOM. Five days afterwards, mice were treated with the following regimen: three cycles of 2.0% DSS in drinking water for 5 days, followed by regular water for 14 days to induce chronic colitis. For antibody treatment, AA98 or mIgG was injected i.p. at 10 mg/kg body weight every other day for three times during each DSS feeding period.

Histopathological and Immunohistochemical Analysis of Mouse Colon Sections

After sacrifice, the colon was removed and cleaned. A 0.5-cm section of the distal third of the colon was collected and stored at −80°C for mRNA isolation. The rest of the colon was rolled as a Swiss roll,34 fixed in 10% formalin neutral buffer solution. Paraffin-embedded sections were cut at 5 μm and stained with H&E solution. Histological scores were evaluated by two independent investigators (H.D. and D.L.), blinded to the source of treatment, as follows33: Epithelium scores 0, normal morphology; 1, loss of goblet cells; 2, loss of goblet cells in large areas; 3, loss of crypts; 4, loss of crypts in large areas. Infiltration scores: 0, no infiltrate; 1, infiltrate around crypt basis; 2, infiltrate reaching to L. mucosae; 3, extensive infiltration reaching the muscularis mucosae and thickening of the mucosa; 4, infiltration of the L. submucosa. The total histological score represents the sum of the epithelium and infiltration score, and ranges from 0 to 8. These sections were further analyzed by immunohistochemistry or immunofluorescence analysis for CD146, CD31 or CD45, CD4. Leukocyte infiltration and blood vessel density were quantified in at least five random areas per section, as previously reported.35

Isolation and Subsequent Analysis of Murine Lamina Propria Lymphocyte

According to the method of Weigmann et al.,36 mouse colons were removed and cleaned. Colons were first predigested with 5 mmol/L EDTA and 1 mmol/L dithiothreitol. Intraepithelial lymphocytes were discarded. The colons were then digested with 0.05 g of collagenase D (Roche, Basel, Switzerland), 0.05 g of DNase I (Sigma-Aldrich). Lamina propria mononuclear cells were isolated on a 40/80 Percoll gradient. Lamina propria mononuclear cells were then stimulated with 1 ng/mL phorbol-12-myristate-13-acetate and 500 ng/mL...
ionomycin (Sigma-Aldrich), harvested and stained with anti–CD3-Percp cy5.5, anti–CD4-APC, anti–IL17A-FITC, or anti–IFN-γ–PE (e-Bioscience). Cells were analyzed using BD LSRRFortessa (BD Biosciences).

Real-Time RT-PCR for Analyzing mRNA Expression Levels

Total RNA was extracted from 30 mg of colon tissues with Trizol reagent (Invitrogen). RNA (2 μg) was reverse transcribed into cDNA by random primers. Real-time PCR analysis was performed on a Corbett 6200 using SYBR Green PCR mix (Toyobo Co., Osaka, Japan). Threshold cycle (Ct) values of GAPDH were subtracted from Ct values of the genes of interest (ΔCt). ΔCt values of mice under different treatments were compared with ΔCt values of untreated mice (naïve). The primers for cytokines were purchased from Qiagen (Quanti-Tect Primer Assays; Valencia, CA), and primers for mouse CD146 and GAPDH were synthesized by Invitrogen using the following sequence (sense; antisense): mCD146: 5′-TCAATGGTTCCGCAACTGATGGA-3′; 5′-GGGAGTTCGGACGTCTGCACC-3′; mGAPDH: 5′-ACTCAAGATTGTCAGCA-3′; 5′-CTCATTCAAGATTGTCAGCA-3′; 5′-GTCTTCTGGTGGCAGTGAT-3′.

ChIP Assays

HUVEC cells were treated with 40 ng/mL TNF-α for 12 hours and fixed for 10 minutes in 1% formaldehyde at room temperature. Chromatin was then sheared to 300 to 1000 bp in length by sonication. Immunoprecipitation was performed with a rabbit anti-p65 polyclonal antibody or control rabbit IgG. Recovered DNA was quantified by PCR and normalized with input DNA. The primers for putative NF-kB binding sites were: NF-621 sense, 5′-GACCTGTGACGGTCTCGC-3′; and antisense, 5′-GGTCTGGCAGTGACGTCTGACC-3′; and for NF-420 sense, 5′-GCCGCAGCGCAGGTCAAG-3′ and antisense, 5′-GTGACAGCGGATGCTGTCGG-3′. GAPDH sense, 5′-CGGATCCACGGTGGTGCTTG-3′; antisense, 5′-TCTCTCAATGGTTGGTGGAGA-3′ was used as control. The PCR products were separated on 1.5% agarose gels.

Dural Luciferase Reporter Assay

A 2-kb CD146 promoter region was amplified from genomic DNA of HUVEC cells and cloned upstream of firefly luciferase in the pGL3 plasmid (Promega, Madison, WI). The mutants of putative NF-kB binding sites of CD146 promoter regions were generated by PCR and confirmed by sequencing. The pGL3 vectors containing WT or mutant CD146 promoter were then transfected into HUVECs, together with pRLTK containing the Renilla luciferase reporter gene and the control empty plasmid. Twelve hours after transfection, cells were treated with 40 ng/mL TNF-α for 24 hours. Firefly and Renilla luciferase activities were then measured with the Dual-Luc Assay Kit (Promega).

Tube Formation Assay

Five experimental groups of HUVECs were prepared with different treatments: mock group (transfected with GFP siRNA and stimulated with PBS, adding 50 μg/mL mIgG right before performing the following experiments), TNF-α stimulation group (transfected with GFP siRNA and stimulated with TNF-α, adding mIgG), restoration group (transfected with CD146 siRNA and stimulated with TNF-α, adding mIgG), blockage group (blockage of NF-kB pathway by BAY11-7082 and stimulated with TNF-α, adding mIgG), and functional antibody AA98 block group (transfected with GFP siRNA and stimulated with TNF-α, adding 50 μg/mL AA98). The relative ratio of TNF-α and CD146 siRNA concentration was adjusted to restore CD146 to its background level of untreated HUVECs as determined by Western blot. Matrigel (BD Biosciences) was thawed at 4°C overnight, and each well of a pre-chilled 96-well plate was coated with 50 μL of Matrigel and incubated at 37°C for 2 hours. As described above, the differentially treated HUVECs were harvested, washed, and then suspended in RPMI 1640 medium containing 10% fetal bovine serum, then added into Matrigel-coated wells. After 12 hours of incubation at 37°C, the cells were observed under a confocal laser scanning microscope (FV500; Olympus, Tokyo, Japan). Endothelial cell tube formation was assessed by endothelial cord branch points and vessel length.

Leukocyte Adhesion and Transendothelium Migration Assays

As described above, HUVECs of different treatment were seeded into a 96-well plate or the upper chamber of transwell plates (96-well insert; pore size 8 μm, Corning Costar; Corning, Corning, NY) for 10,000 cells per well. Subsequently, cells were incubated for 24 hours, stimulated by 40 ng/mL TNF-α or PBS. Thirty thousand fresh isolated peripheral mononuclear cells (PBMCs) or THP-1 cells/well were stained with carboxyfluorescein succinimidyl ester and added to the plate or upper chamber. For the adhesion assay, after incubation for 1 hour, wells were washed three times with 0.2 mL of warmed assay medium to remove non-adherent leukocytes. The adhered leukocyte were harvested and analyzed by flow cytometry. For the transmigration assay, the leukocytes were allowed to migrate through the endothelial monolayer for 12 hours. The transmigrated cells in lower chamber were collected and analyzed by flow cytometry.

In Vitro T-Cell Differentiation

CD4+ T cells isolated using a CD4+ T-cell Isolation Kit II (Miltenyi Biotec, Bergisch Gladbach, Germany) from
spleen and lymph nodes were activated with plate-bound 3 μg/mL anti-CD3 (145-2C11; BD Biosciences) and 3 μg/mL CD28 (37.51; BD Biosciences). The following Th17 culture condition was used: 0.5 ng/mL TGF-β (R&D Systems, Minneapolis, MI), 10 ng/mL IL-6 (R&D Systems), 10 ng/mL IL-1β (R&D Systems), 10 ng/mL IL-21 (R&D Systems), 10 μg/mL anti–IFN-γ (XMG1.2; BD Biosciences), 10 μg/mL anti–IL-12p40, and 10 μg/mL anti–IL-4 (11B.11; National Cancer Institute, Frederick, MD). The following regulatory T cell (Treg) culture condition was used: 5 ng/mL TGF-β, 50 U/mL IL-2 (National Cancer Institute), 10 μg/mL anti–IL-4, 10 μg/mL anti–IL-12p40, 10 μg/mL anti–IFN-γ. Th0 cells were maintained with 50 U/mL IL-2. These T cells were activated for 4 days before being used for transmigration assays.

Statistical Analysis

The results are presented as means ± SEM. Differences between groups were calculated for significance by one-way or two-way analysis of variance with a Tukey test using GraphPad Prism software version 5 (GraphPad Software, La Jolla, CA). For statistical analysis of DAI, two-tailed paired Student’s t-test was used. A P value <0.05 was considered significant. All of the experiments were repeated at least three times independently, and representative data were shown.

Results

Colitis Severity Is Reduced in Endothelial CD146 Knockout Mice

It has been reported that CD146 expression is increased on blood vessels of IBD patients by an unknown mechanism.23,24 Consistently, we found both increased endothelial CD146 expression and enhanced angiogenesis in colon tissues from DSS-induced colitis model mice; however, no epithelial CD146 expression was observed (Figure 1A). To investigate the function of endothelial CD146 in inflammation during colitis, we compared the susceptibility of the conditional endothelial cell CD146 knockout mice (CD146EC-KO) and WT mice to DSS-induced acute colitis. In the CD146EC-KO mice, we confirmed the depletion of CD146 in vascular endothelial cells (Supplemental Figure S1) but not in other CD146-expressing cells such as hematopoietic stem cells (CD117+CD146+) and CD146+ lymphocytes, as we reported previously.31 In addition, no significant differences in colon histology, colon cytokine mRNA levels, and lymphocyte subpopulations were observed between naïve (nontreated) CD146EC-KO mice and WT mice (Supplemental Figure S2). When DSS induction was performed, deletion of CD146 in endothelial cells markedly dampened disease severity, including weight loss, occult and rectal bleeding, and diarrhea of mice, as shown by the DAI in Figure 1B.

Necropsy of mice at day 9 revealed alleviated edema and shortening of the colons in DSS-fed CD146EC-KO mice compared with WT mice (data not shown). In addition, less severe epithelial damage and lymphocyte infiltration were found in DSS-induced CD146EC-KO mice by histological examination of colonic sections (Figure 1B).

Using CD31 and CD45 marking blood vessel endothelium and pan-lymphocytic cells, respectively, we noted decreases in both vessel density and leukocyte infiltration in colon tissue of CD146EC-KO mice compared with WT mice (Figure 1, C and D). Because Th1 and Th17 are two major proinflammatory lymphocytes that have been implicated in IBD,37 we investigated the possibility that endothelial CD146 was involved in recruiting these lymphocytes to sites of active inflammation. The infiltrated lymphocytes in the lamina propria of DSS-treated CD146EC-KO and WT mice were isolated and analyzed. We found fewer CD4+ lymphocytes and a lower percentage of Th1 and Th17 cells present in the colon of CD146EC-KO mice than that of WT mice (Figure 1E). Also, the IFN-γ IL17+ double-positive cells were reduced (Supplemental Figure S3). In addition, mRNA levels of cytokines in the colon tissue were measured to evaluate the inflammation status during colitis. The expression of major proinflammatory cytokines, including TNF-α, IL1-β, and IL6, were all reduced in CD146EC-KO mice (Figure 1F).

Furthermore, a TNBS acid–induced colitis model was used to confirm the possible role of endothelial CD146 in mediating lymphocyte infiltration, because TNBS-induced colitis was characterized by a predominant Th1-mediated immune response.38,39 Colitis severity was significantly ameliorated in TNBS-induced CD146EC-KO mice, compared with induced WT mice (Supplemental Figure S4). The infiltration of CD4+ lymphocyte, as detected by immunofluorescence and flow cytometry, was markedly reduced in the CD146EC-KO model mice. We also found reduced infiltration of Th1 and Th17 cells in the colons of CD146EC-KO model mice, compared with that of WT model mice.

Taken together, these data indicate that endothelial CD146 plays a crucial role in the pathogenesis of colitis by promoting inflammation.

Eliminating CD146 Inhibits Colitis-Associated Colorectal Carcinogenesis

Chronic inflammation contributes to carcinogenesis. To determine whether the role of CD146 in inflammation is of importance in CAC, we evaluated the susceptibility of WT and CD146EC-KO mice to AOM-DSS–induced CAC (Figure 2A). During the course of DSS treatment, profound body weight loss and bloody diarrhea were observed in WT mice, compared with much milder symptoms in CD146EC-KO mice (data not shown). When mice were sacrificed at day 62, reduced tumor incidence and smaller tumor volume were observed in CD146EC-KO mice compared with WT mice (Figure 2, B–D). Immunofluorescence analysis revealed lower blood vessel density and a decreased level of CD4+
leukocyte infiltration in the tumor regions of CD146EC-KO mice (Figure 2, E and F). Moreover, significantly decreased mRNA level of TNF-α, IL1-β, IFN-γ, and IL6 was found in colon samples from CD146EC-KO mice compared with those from treated WT mice (Figure 2G). Also, COX-2, whose expression is pivotal in CAC,40 was significantly reduced in CD146EC-KO mice compared to WT mice. Consistently, by analyzing the cytokine concentration present in serum, we
observed decreased TNF-α, IFN-γ, and IL6 levels in CD146EC-KO mice (data not shown). As a control, no significant differences in colon histology, colon cytokine mRNA levels, and lymphocyte subpopulations were observed between naive CD146EC-KO mice and WT mice used in the CAC model (Supplemental Figure S5). We next analyzed whether CD146 contributes to the transformation of epithelial cells in a cell autonomous manner. Immunohistochemistry analysis showed no CD146 expression in epithelial cells, which was consistent with the expression pattern of CD146 in human colorectal cancer (Supplemental Figure S6A). Moreover, β-catenin translocation was analyzed because its overexpression and translocation are important events in colorectal carcinogenesis. In the tumor region of both WT and CD146EC-KO mice, overexpression and translocation of β-catenin from membrane to cytoplasm and nucleus were observed. However, no significant difference in β-catenin expression pattern was found between WT and CD146EC-KO mice (Supplemental Figure S6B). Meanwhile, immunofluorescence analysis of Ki-67 and TUNEL assay revealed no significant differences in proliferation and apoptosis (Supplemental Figure S6, C and D). These data together suggest that endothelial CD146 facilitates carcinogenesis mainly by promoting inflammation.

CD146 Is Up-Regulated by TNF-α and IL1-β through NF-κB Transactivation

To elucidate the underlying mechanism of endothelial CD146 involvement in colitis and CAC, we examined the expression of CD146 after treating HUVECs and sEnd.1 with different cytokines. We found that TNF-α and IL1-β, but not IFN-γ, IL6, or VEGF, could up-regulate CD146.
expression in HUVEC cells (data not shown). TNF-α up-regulated CD146 in a dose- and time-dependent manner, at both the protein and mRNA levels (Figure 3, A–C). Similarly, this phenomenon was observed in HUVECs with IL1-β treatment (Supplemental Figure S7, A and B) and mouse endothelioma End.1 cells with mTNF-α treatment (Supplemental Figure S7C).

Because both TNF-α and IL1-β activate the NF-κB pathway, we suspected that CD146 up-regulation might be mediated by NF-κB. Indeed, we found that the NF-κB pathway inhibitor BAY11-7082 blocked CD146 up-regulation, whereas the inhibitors of other pathways did not (Figure 3D). In addition, knockdown of NF-κB subunit p65 also abolished the up-regulation of CD146 by TNF-α (Supplemental Figure S8A). These results indicate that TNF-α induces CD146 up-regulation mainly through NF-κB pathway.

To determine the mechanism of NF-κB–mediated CD146 up-regulation, we analyzed the promoter region of CD146 (MCAM) using the online database ECR Browser and found two putative NF-κB binding sites (NF-κB site 1 at −621 and NF-κB site 2 at −420). Using a luciferase reporter gene driven by the CD146 promoter region containing the putative NF-κB binding sites, we found that TNF-α robustly enhanced the luciferase activity, whereas mutations of the two putative sites abolished such activity (Figure 3E). This result indicates that both the putative NF-κB binding sites are vital for TNF-α–induced CD146 expression. To further examine whether NF-κB directly bound to these putative sites, a ChIP assay was performed against the p65 subunit of NF-κB. A p65-specific polyclonal antibody enriched the CD146 promoter region in a TNF-α dependent fashion (Figure 3F). These results indicate that the NF-κB p65 subunit binds to the sites directly. Taken together, these results suggest that after stimulation, NF-κB directly binds to the CD146 promoter region and modulates CD146 expression.

Enhanced Endothelial CD146 Expression Promotes Angiogenesis and Endothelium–Leukocyte Interaction

To determine the effect of the up-regulation of CD146 expression on modulating endothelial functions, we modulated the expression of CD146 on HUVECs with TNF-α...
Enhanced endothelial CD146 expression promotes endothelial cell tube formation and endothelium–leukocyte interaction. A: CD146 expression was analyzed by Western blot in HUVECs with indicated treatments, before the tube formation and endothelium–leukocyte interaction assays. GFP siRNA and mIgG were used as control siRNA and antibody, respectively. B: Representative images of tube-formation assay of HUVEC with indicated treatments. The bar graph shows the quantification as relative tube length with the GFP siRNA+mIgG group set as 100%. C: Adhesion assay of fresh isolated PBMCs from a healthy donor adhered to HUVECs with indicated treatments. Transendothelial migration assay of fresh isolated PBMCs (sEnd.1) with the indicated treatments. D: Analysis of in vitro differentiated murine Th1, Th17, Treg, and Th0 lymphocyte transmigration across the endothelial monolayer formed by mouse endothelioma cells (THP-1) with the indicated treatments. E: Analysis of in vitro differentiated murine Th1, Th17, Treg, and Th0 lymphocyte transmigration across the endothelial monolayer formed by mouse endothelioma cells (sEnd.1) with the indicated treatments. *P < 0.05, **P < 0.01, ***P < 0.001 (B–E). |P < 0.05, |P < 0.01 versus the GFP siRNA+mIgG group (F). n = 3 (B–F). Original magnification, ×100 (B).

stimulation, and then examined the effect of such treatment first on tube formation. The relative ratio of TNF-α and CD146 siRNA concentration was adjusted to restore CD146 to the baseline level of untreated HUVECs as determined by Western blot (Figure 4A). TNF-α stimulation strongly promoted tube formation of HUVECs on Matrigel (Figure 4B), which appeared to be mediated by CD146 because CD146-specific siRNA, NF-κB inhibitor BAY11-7082, or treatment with the anti-CD146 functional antibody AA98 all abolished this effect (Figure 4B and Supplemental Figure S9A). A similar CD146 dependence was observed in TNF-α–induced HUVEC migration (data not shown). These data were further confirmed by using a second CD146 siRNA or p65 siRNA (Supplemental Figure S8, B–E). Taken together, these results indicate that enhanced CD146 expression induced by TNF-α promotes angiogenesis.

Furthermore, we analyzed the effect of CD146 on the interaction between endothelium and inflammatory cells. Enhanced endothelial CD146 expression promoted PBMC adhesion to and transmigration across endothelial monolayer (Figure 4, C and D, and Supplemental Figures S6F and S9B). A similar effect was observed when using monocyte THP-1 cells (Figure 4E). Because increased Th1 and Th17 lymphocyte infiltration was observed in the lamina propria from WT mice compared to CD146EC-KO mice, we suspected that CD146 on endothelial cells might facilitate recruitment of these proinflammatory lymphocytes. To test this possibility, a transendothelial migration assay was performed with in vitro differentiated Th1, Th17, Treg, and Th0 cells. Interestingly, endothelial cells with TNF-α–induced CD146 expression recruited more Th1 and Th17 lymphocytes than Treg and Th0 cells, which could be blocked by CD146 knockdown, NF-κB inhibition, or antibody AA98 (Figure 4F), suggesting that enhanced expression of endothelial CD146 facilitates proinflammatory Th1 and Th17 infiltration.

CD146 has been reported to be expressed in some subsets of leukocytes. We first tested whether AA98 affects the function of CD146− and CD146+ lymphocytes. Murine CD146− and CD146+ lymphocytes isolated by flow cytometry were treated with AA98 or mIgG for 48 hours before they were used to transmigrate across the sEnd.1 endothelial monolayer. CD146+ lymphocytes were observed to have a greater transmigrating ability than CD146− lymphocytes. However, AA98 did not affect the transmigration of CD146+ lymphocyte (Supplemental Figure S10). Also, AA98 had no effect on differentiation and proliferation of lymphocyte in vitro as we previously reported.11 We next examined whether AA98 affects the function of macrophage. We found that AA98 did not affect LPS-induced NF-κB activation and cytokine production in peritoneal exudate cells (Supplemental Figure S11, A–C). In addition, treating peritoneal exudate cells with AA98 did not affect its migration (Supplemental Figure S11D). Similar results were found when we treated bone marrow–derived macrophages with AA98 (data not shown). Therefore, although CD146 is expressed on subsets of lymphocytes and macrophages, our data show that AA98 does not affect their function. All together, these data indicate that enhanced endothelial CD146 expression facilitates angiogenesis, as well as...
endothelium–leukocyte interactions, whereas anti-CD146 antibody AA98 predominantly targets endothelial CD146, rather than CD146 on leukocytes, to function.

**Targeting CD146 by Antibody AA98 Attenuates Colitis and Prevents CAC in Murine Models**

Because CD146 plays an important role in colitis pathogenesis and AA98 blocks CD146 function in *in vitro* experiments, we examined the therapeutic value of anti-CD146 antibody AA98 in mouse models. By administered AA98 or an isotypic control (mlG) preventively to the DSS-treated mice (during the DSS induction process), we found that AA98 significantly alleviated disease severity (Figure 5A). To mimic the long-lasting and relapsing property of IBD, we established a chronic colitis murine model, and administrated AA98 after the onset of disease. Because IBD is a clinically heterogeneous disease with complex mechanisms, a combination of drugs targeting distinct biomarkers might be considered a potential approach. Thus, we also tested the combination treatment of anti-CD146 antibody AA98 and anti–TNF-α antibody V1q in this chronic colitis murine model. DAI and histological score were dramatically reduced in the AA98 treatment group and the V1q treatment group, especially the combination treatment (AA98+V1q) group, compared to the mlG treatment group (Figure 5B). Consistent with the observation in CD146EC-Ko colitis model mice, we found a decrease in blood vessels and lymphocytic cell infiltration, both in AA98-treated and V1q-treated mice, whereas a more significant decrease was observed in the combination treatment group (Supplemental Figure S12, A and B). In addition, the expression levels of proinflammatory cytokines were also reduced (Supplemental Figure S12C). These data indicate that AA98, especially in combination with anti–TNF-α antibody, could be an efficient treatment for colitis.

Furthermore, because AA98 treatment appeared to ameliorate the inflammatory response in colitis, we tested whether it could protect mice with colitis from developing tumors by treating CAC mice with either mlG or AA98 during the DSS induction process. Interestingly, tumor incidence and tumor volume were drastically reduced in AA98-treated mice (Figure 5C). Collectively, these data indicate the therapeutic value of anti-CD146 antibody AA98 in treating colitis and preventing CAC.

**Discussion**

CD146 has been reported to be associated with inflammation diseases through unknown mechanisms. In this study, we show that enhanced CD146 expression on endothelial cells facilitates angiogenesis and leukocyte extravasation, suggesting that endothelial CD146 may serve as a critical molecular promoting inflammation and linking chronic colitis and CAC. Notably, targeting CD146 by either antibody blocking or the use of a conditional knockout significantly alleviated inflammation during acute and chronic colitis, and also decreased tumor incidence and tumor volume in a murine
CAC model. This is the first evidence that the modulation of inflammation and angiogenesis simultaneously by targeting a single adhesion molecule on endothelial cells can prevent CAC.

Our previous study established CD146 as a signaling receptor, more interestingly, a VEGF coreceptor on the surface of endothelial cells, capable of activating the p38/IKK/NF-κB signaling cascade, inducing endothelial activation and cytoskeleton remodeling, leading to tumor angiogenesis. However, the regulatory mechanisms of CD146 and its function in inflammation remain unknown. Here, we study the upstream regulation mechanism of CD146 and present evidence that endothelial CD146 expression is induced directly by NF-κB transactivation stimulated by TNF-α. In turn, CD146 overexpression on endothelial cells promotes angiogenesis, as well as facilitates recruitment of proinflammatory Th1 and Th17 lymphocytes to the sites of inflammation. Such a dual function of endothelial CD146 could exacerbate the inflammation in the affected tissues, resulting in a pathological cycle leading the initial inflammation to chronic inflammation and at last to carcinogenesis. These results explained the function and regulation mechanism of enhanced CD146 expression on endothelial cells during IBD, which were previously observed by Tsolakidou et al. and Bardin et al. in intestinal biopsies from IBD patients. In addition, it further provides the evidence that modulation of the inflammatory environment by targeting vascular endothelial cells could reduce the incidence of inflammation-associated carcinogenesis.

Here, we focused on the function of CD146 in endothelial cells. However, CD146 expression has also been reported on Th17 lymphocytes and certain macrophage subpopulations. As shown in our recent publication, the knockout of endothelial CD146 in CD146EC-KO mice does not impair CD146+ lymphocytes. Furthermore, the anti-CD146 antibody AA98 does not affect the proliferation or activation of either CD146+ or CD146- lymphocytes. Interestingly, we found that pretreating CD146+ lymphocytes alone with AA98 did not affect their ability of transendothelial migration, whereas pretreating endothelial cells alone did block leukocyte transmigration. In addition, we found that AA98 did not affect LPS-induced NF-κB activation and cytokine production in peritoneal exudate or bone marrow-derived macrophages. Also, treating macrophages with AA98 did not block their migration. Although the role of CD146 in leukocytes remains to be determined, our data suggest that enhanced endothelial CD146 expression facilitates leukocyte (either CD146+ or CD146−) transendothelial migration. In addition, the AA98 antibody selectively targets endothelial CD146, but not CD146 on lymphocytes or macrophages, to inhibit the extravasation of leukocytes.

Of note, our CD146EC-KO mice were generated using a Tg(Tek-Cre) system. Because Tie2 (Tek) was reportedly expressed on a subset macrophage, we cannot exclude a possible effect of CD146+ macrophages involved in colitis and CAC. However, our data have shown that blocking CD146 by AA98 did not have an effect on activation and migration of macrophages, whereas AA98 did alleviate colitis and CAC. Therefore, we suspect that the CD146 expressed on certain macrophages and lymphocytes may play a minor role compared with endothelial CD146. Detailed analysis of CD146 on macrophages awaits the generation of macrophage-specific CD146 knockout mice.

The current treatment of IBD mainly focuses on inhibition of cytokines. Infliximab, an antibody against TNF-α, was approved by the US Food and Drug Administration for treating UC. Many inhibitors and antibodies against TNF-α (Etanercept, Oncept), IFN-γ (Fontolizumab), IL12p40, and IL6 receptor are now under clinical evaluation. Moreover, side effects such as increased risk of infection and even lymphoma have been indicated. Moreover, clinical trial data suggest that a considerable proportion of patients receiving anti-TNF-α therapy may become refractive in the first few years. Therefore, anti-adhesion molecule therapy could be an alternative therapeutic intervention. In this study, we tested the efficacy of our anti-CD146 antibody AA98 and also a combination treatment of anti-TNF-α and anti-CD146. AA98 markedly attenuated experimental colitis, and the combination of anti-CD146 and anti-TNF-α had an even more significant effect. Histological examination revealed that AA98 treatment preserved a more integral epithelial structure with less leukocyte infiltration, although the change in DAI indicated that targeting TNF-α ameliorated the clinical symptoms more rapidly than AA98 treatment. This may be due to the different mechanisms underlying the action of these two molecules. Thus, the combination therapy of anti-CD146 and anti-TNF-α may have a synergistic effect and hold a promising potential.

In this study, we propose that CD146 plays its role in promoting CAC mainly by facilitating the inflammatory environment, because no CD146 expression was found in epithelial cells of the tumor. It has been now well accepted that decreased inflammation leads to decreased epithelial proliferation and decreased DNA damage, which results in reduced carcinogenesis. Thus, any agent that reduces inflammation in the AOM-DSS model would be expected to reduce CAC, making it impossible to distinguish between the effects of targeting CD146 on colitis activity and a separate antineoplastic effect. However, a unique characteristic of CD146 is that CD146 plays a dual role in endothelial cells—facilitating leukocyte extravasations and angiogenesis simultaneously. Because the development of tumor depends indispensably on angiogenesis, targeting endothelial CD146 not only affects the inflammatory environment, but also prohibits tumor development directly. Nevertheless, further study should be done to fully understand the function and mechanism of CD146 in the course of tumor development.

Taken together, our data demonstrate that endothelial CD146 plays a vital role in inflammation. Its involvement in the recruitment of inflammatory lymphocytes and promotion
of angiogenesis, may serve as a critical linking between colitis and colorectal carcinogenesis. Targeting CD146 may be a therapeutic option in treating colitis, as well as a preventative method for colitis-associated colorectal carcinogenesis.

Supplemental Data

Supplemental material for this article can be found at http://dx.doi.org/10.1016/j.ajpath.2014.01.031.

References


