Therapeutic Role of Rifaximin in Inflammatory Bowel Disease: Clinical Implication of Human Pregnane X Receptor Activation

Jie Cheng, Yatrik M. Shah, Xiaochoao Ma, Xiaoyan Pang, Toshiya Tanaka, Tatsuhiko Kodama, Kristopher W. Krausz, and Frank J. Gonzalez

Laboratory of Metabolism, Center for Cancer Research, National Cancer Institute, National Institutes of Health, Bethesda, Maryland (J.C., X.P., K.W.K., F.J.G.); Departments of Molecular and Integrative Physiology and Internal Medicine, Division of Gastroenterology, University of Michigan School of Medicine, Ann Arbor, Michigan (Y.M.S.); Department of Pharmacology, Toxicology, and Therapeutics, University of Kansas Medical Center, Kansas City, Kansas (X.M.); and Laboratory for Systems Biology and Medicine, University of Tokyo, Tokyo, Japan (T.T., T.K.)

Received May 12, 2010; accepted July 8, 2010

ABSTRACT

Human pregnane X receptor (PXR) has been implicated in the pathogenesis of inflammatory bowel disease (IBD). Rifaximin, a human PXR activator, is in clinical trials for treatment of IBD and has demonstrated efficacy in Crohn’s disease and active ulcerative colitis. In the current study, the protective and therapeutic role of rifaximin in IBD and its respective mechanism were investigated. PXR-humanized (hPXR), wild-type, and Pxr-null mice were treated with rifaximin in the dextran sulfate sodium (DSS)-induced and trinitrobenzene sulfonic acid (TNBS)-induced IBD models to determine the protective function of human PXR activation in IBD. The therapeutic role of rifaximin was further evaluated in DSS-treated hPXR and Pxr-null mice. Results demonstrated that prednimistration of rifaximin ameliorated the clinical hallmarks of colitis in DSS- and TNBS-treated hPXR mice as determined by body weight loss and assessment of diarrhea, rectal bleeding, colon length, and histology. In addition, higher survival rates and recovery from colitis symptoms were observed in hPXR mice, but not in Pxr-null mice, when rifaximin was administered after the onset of symptoms. Nuclear factor κB (NF-κB) target genes were markedly downregulated in hPXR mice by rifaximin treatment. In vitro NF-κB reporter assays demonstrated inhibition of NF-κB activity after rifaximin treatment in colon-derived cell lines expressing hPXR. These findings demonstrated the preventive and therapeutic role of rifaximin on IBD through human PXR-mediated inhibition of the NF-κB signaling cascade, thus suggesting that human PXR may be an effective target for the treatment of IBD.

Introduction

Rifaximin [Rifax; Xifaxan; 4-deoxy-4’-methylpyrido[1’,2’,1,2]imidazo[5,4-c]rifamycin SV], a nonsystemic rifamycin-derived antibiotic that exhibits low gastrointestinal absorption while retaining potent antibacterial activity (Koo and Dupont, 2010), was approved in 2004 for the treatment of traveler’s diarrhea (Laustsen and Wimmett, 2005). Clinical trials have also indicated the potential role for rifaximin in inflammatory bowel disease (IBD) (Day and Gearry, 2010). Rifaximin seems to have more efficacy in the therapy of irritable bowel syndrome than do other antibiotics such as neomycin, doxycycline, amoxicillin/clavulanate, and ciprofloxacin (Yang et al., 2008). Other studies revealed that two-thirds of adult patients with Crohn’s disease entered remission after rifaximin therapy (Shafran and Burgunder, 2010). A retrospective review revealed a well tolerated and favorable role for rifaximin in pediatric IBD (Muniyappa et al., 2009; Trehan et al., 2009). However, the role of rifaximin in irritable bowel syndrome and IBD therapy and its mechanisms of action are not understood.

A previous study demonstrated that rifaximin is a gut-
specific human pregnane X receptor (PXR) agonist (Ma et al., 2007b). PXR is a ligand-activated transcription factor important for its induction of drug transport and metabolism, in particular induction of the cytochrome P450 CYP3A4 involved in the metabolism of many clinically used drugs. Recent data indicate that PXR may be involved in IBD. Gene expression analysis of colon tissues from ulcerative colitis and patients with Crohn’s disease demonstrated a significant inhibition of PXR and its target genes compared with normal intestinal samples (Langmann et al., 2004). Specific polymorphisms in the PXR locus that are associated with a decrease in PXR activity are correlated with an increased susceptibility to IBD (Dring et al., 2006). PXR inhibits the proinflammatory transcription factor NF-κB, providing a potential molecular mechanism that links PXR signaling and inflammation (Xie and Tian, 2006; Zhou et al., 2006a). For example, activation of mouse PXR ameliorates dextran sulfate sodium (DSS)-induced colitis via inhibition of NF-κB (Shah et al., 2007). However, PXR ligands are structurally diverse and as noted above display species specificity (Ma et al., 2008). Rifaxipicin [3-(4-methylpiperazinyliminomethyl) rifamycin SV] is an agonist for human PXR but does not activate rodent PXR, whereas pregnenolone-16α-carotinol, a rodent-specific PXR agonist, does not activate human PXR (Zhou et al., 2009). Because of the specific activation of rifaximin on human PXR, the role of this drug in therapy of IBD has been a focus of interest.

In the current study, the preventive and therapeutic role of rifaximin was assessed in DSS- or TNBS-induced IBD models. Rifaxipicin (Ma et al., 2007b), as another rifamycin-derived antibiotic and human PXR agonist (Bertilsson et al., 1998), was compared with rifaximin. Because rodent models cannot accurately predict inducers and potential drug–drug interactions mediated by human PXR due to different responses to PXR ligands (Jones et al., 2000), PXR-humanized (hPXR) mice were used to investigate the potential role of human PXR activation in the mouse IBD models. The hPXR mice express human PXR in the Pxr-null background and respond only to human PXR-specific ligands (Ma et al., 2007a). Wild-type (WT) and Pxr-null mouse lines were also assessed to distinguish the antibiotic role from its PXR ligand effect. Rifaxipin treatment demonstrated no protection against mouse IBD and exacerbated the severity of IBD. However, the hPXR mice treated with rifaximin demonstrated significant protection and a potential therapeutic role of human PXR in IBD. No protection of rifaximin was observed in wild-type and Pxr-null mice, thus demonstrating that rifaximin functions through activating human PXR and not through its gut-specific antibiotic effects. In addition, rifaximin inhibited the NF-κB signaling cascade in a human PXR-dependent manner. These data provide the first mechanistic evidence by which rifaximin and human PXR influence experimental IBD.

**Materials and Methods**

**Animals.** hPXR, wild-type, and Pxr-null male mice were housed in temperature- and light-controlled rooms and given water and pelleted chow ad libitum. The hPXR mice express the human PXR in the Pxr-null background. All animal experiments were carried out in accordance with the Institute of Laboratory Animal Resources guidelines and approved by the National Cancer Institute Animal Care and Use Committee.

**Experimental Design.** Two- to 3-month-old hPXR, wild-type, and Pxr-null male mice were subjected to DSS- and TNBS-induced IBD models. Mice were placed into four groups (n ≥ 6 per group) in the DSS-induced IBD study: control, DSS, rifaximin or rifampicin, and DSS treatment after rifaximin or rifampicin pretreatment. The control and DSS groups received a control diet for 11 consecutive days, and mice in the rifaximin (Salix Pharmaceuticals, Inc., Morrisville, NC) or rifampicin (Sigma-Aldrich, St. Louis, MO) groups received rifaximin (1 mg/kg/day) or rifampicin (3 mg/kg/day or 10 mg/kg/day), respectively, in the diet for 11 consecutive days. Dosages of rifaximin or rifampicin were calculated based on mouse daily dietary intake. Mice in the rifaximin and DSS or rifampicin and DSS groups received the compounds in the diet for 11 days, and on the 5th day of treatment the mice were administered 2.5% DSS (MP Biomedicals, Solon, OH) in drinking water (wt/v), and all mice were killed 11 days after treatment with the rifaximin or rifampicin diets or the control diet. Serum samples were collected by retro-orbital bleeding. hPXR and Pxr-null mice were used for the therapeutic studies (n ≥ 10 per group) in which mice were treated with 2.5% DSS and control diet for 5 consecutive days, then DSS was withdrawn and rifaximin was administered and compared with vehicle-treated mice. The mice were observed for 7 consecutive days after the onset of rifaximin treatment. The TNBS-induced IBD protocol was also used to confirm the role of rifaximin and human PXR in IBD (n ≥ 6 per group). A 2-mg aliquot of TNBS in 50% ethanol was administered intrarectally to anesthetized mice via a round-bottom (50.8 mm) needle (Pupper, Lake Success, NY), followed by control diet or rifaximin diet treatment for 6 consecutive days. The mice were killed 5 days after TNBS administration, and the colons were flushed and resected. All of the above IBD studies were repeated twice. All mice were killed by CO2 asphyxiation at the end of the study, and tissue samples were harvested and stored at −80°C before analysis.

**Colitis Evaluation.** Daily changes of body weight, diarrhea, rectal bleeding, and bloody stool were assessed and reported as a score from 0 to 4. For assessment of macroscopic colon damage, the colon was opened longitudinally, flushed with phosphate-buffered saline, and fixed in 10% buffered formalin. Colitis was measured by blinded analysis on a routine hematoxylin and eosin (H&E)-stained section according to the morphological criteria described previously (Cooper et al., 1993).

**RNA Analysis.** RNA was isolated and mRNA expression was assessed by quantitative real-time PCR (qPCR) as described previously (Cheng et al., 2009).

**Cell Lines and Luciferase Assays.** Human epithelial colorectal adenocarcinoma cells (Caco-2) and human colonic epithelial cells (HT-29) were grown at 37°C with 5% CO2 in Dulbecco’s modified Eagle’s medium (Invitrogen, Carlsbad, CA) supplemented with 10% fetal bovine serum (Gemini Bio-Products, Woodland, CA) and 1% penicillin/streptomycin (Invitrogen). Caco-2 and HT-29 cells were seeded at a density of 5 × 104 cells/well in 24-well plates. Expression vectors for human PXR, retinoid X receptor (RXR), and NF-κB luciferase reporters were transfected into cells by using the FuGene transfection reagent (Roche Diagnostics, Indianapolis, IN). Twenty-four hours after transfection, the cells were incubated with DMSO (vehicle) or rifaximin for 24 h, followed by incubation with tumor necrosis factor α (TNFα) (10 ng/ml) for 24 h. A standard dual luciferase assay was used and normalized to a cotransfected control reporter plasmid (Promega, Madison, WI).

**Protein Analysis of Human PXR and Microsomal Stearoyl-CoA Desaturase-1.** Fresh colon tissues were collected, and epithelial cells isolated. Nuclear protein was isolated with a NE-PER kit (Thermo Fisher Scientific, Rockford, IL). Liver tissues were collected and homogenized in ice-cold buffer (1.15% KCl, 50 mM Tris-HCl, and 1 mM EDTA, pH 7.4) and microsomes were prepared by differential centrifugation. Pellets were resuspended in 100 mM Tris-HCl (pH 7.4), 0.1 mM EDTA, 0.1 mM dithiothreitol, 0.15 M KCl, and 20% (v/v) glycerol, aliquoted, and stored at −80°C. Primary antibodies to human PXR (H5017) and stearoyl-CoA desaturase-1 (SCD-1) (Santa
Cruz Biotechnology, Inc., Santa Cruz, CA) diluted 1:1000 with Tris-buffered saline/Tween 20, followed by peroxidase-conjugated antimouse IgG diluted 1:10,000 with Tris-buffered saline/Tween 20, were used for Western blot analysis. Nuclear hepatocyte nuclear factor 4α (HNF4α) and microsomal glycerol 3-phosphate dehydrogenase were used as loading controls.

**Metabolomics Analysis of Serum.** A 1-μl aliquot of diluted serum samples was injected into a Waters (Milford, MA) UPLC-TOFMS system. An Acquity UPLC BEH C18 column (Waters) was used to separate chemical components at 35°C. The mobile-phase flow rate was 0.5 mL/min with an aqueous acetonitrile gradient containing 0.1% formic acid over a 10-min run (0% acetonitrile for 0.5 min to 20% acetonitrile by 5 min to 95% acetonitrile by 9 min, then equilibration at 100% water for 1 min before the next injection). The QTOF Premier mass spectrometer (Waters) was operated in the positive electrospray ionization mode. Capillary voltage and cone voltage were maintained at 3 kV and 20 V, respectively. Source temperature and desolvation temperature were set at 120 and 350°C, respectively. Nitrogen was used as collision gas. For accurate mass measurement, the time of flight mass spectrometry was calibrated with sodium formate used as collision gas. The QTOF Premier mass spectrometer (Waters) was operated in the positive electrospray ionization mode. Capillary voltage and cone voltage were maintained at 3 kV and 20 V, respectively. Source temperature and desolvation temperature were set at 120 and 350°C, respectively. Nitrogen was used as collision gas. For accurate mass measurement, the time of flight mass spectrometry was calibrated with sodium formate solution (range m/z 100-1000) and monitored by the intermittent injection of the lock mass sulfadimethoxine ([M + H]^+ = 311.0814 m/z) in real time. Mass chromatograms and mass spectral data were acquired and processed by MassLynx software (Waters) in centroid format.

**Principal Components Analysis of Serum Metabolomic Data.** Chromatographic and spectral data were deconvoluted by MarkerLynx software (Waters). A multivariate data matrix containing information on sample identity, ion identity (retention time and m/z), and ion abundance was generated through centroiding, deisotoping, filtering, peak recognition, and integration. The intensity of each ion was calculated by normalizing the single ion counts versus the total ion counts in the whole chromatogram. The data matrix was further exported into SIMCA-P software (Umetrics, Kinnelon, NJ) and transformed by mean-centering and Pareto scaling, a technique that increases the importance of low abundance ions without significant amplification of noise. Principal components of serum were generated by principal components analysis to represent the major latent variables in the data matrix and were described in a scores scatter plot.

**Data Analysis.** Experimental values are expressed as mean ± S.D. Statistical analysis was performed with two-tailed Student’s t tests, and percentage of survival was analyzed by Prism 5.0 (GraphPad Software, Inc., San Diego, CA) with survival curve statistical analysis. A p value of < 0.05 was considered statistically significant.

**Results**

**Rifampicin Does Not Protect hPXRx Mice in the DSS-Induced Colitis Model.** Rifampicin is a derivative antibiotic of rifamycin SV, which is a systemic drug widely studied as a typical human PXR agonist. Long-term treatment in tuberculosis therapy is thought to increase hepatotoxicity (Tostmann et al., 2008). Here, rifampicin was investigated for the treatment of DSS-induced colitis. Effective dosages of 3 mg/kg/day (Xie et al., 2000) and 10 mg/kg/day were selected (Cheng et al., 2009). hPXRx mice pretreated with rifampicin followed by DSS treatment demonstrated no protection against DSS-induced colitis compared with DSS treatment alone. No significant changes were observed in control and rifampicin-treated (3 or 10 mg/kg/day) mice in the absence of DSS. It is noteworthy that high doses of rifampicin, combined with DSS, led to more severe colitis compared with coadministration of low dosage of rifampicin with DSS as revealed by decreased body weight and increased diarrhea and rectal bleeding scores (Fig. 1). Expression of hepatic SCD-1, a lipogenic enzyme regulated by PXR (Zhou et al., 2006b), which is protective in IBD (Chen et al., 2008), was repressed at the mRNA and protein levels after DSS alone. Whereas 3 mg/kg/day rifampicin treatment combined with DSS did not alter these low levels, expression was markedly reduced with 10 mg/kg/day of the drug (Supplemental Fig. 1). This reduction of SCD-1 was associated with a reduction of unsaturated lysophosphatidylcholine in plasma. Human PXR target genes in liver and colon were induced and proinflammatory factors were unchanged by rifampicin in the DSS IBD model (Supplemental Fig. 2).

**Rifaximin Protects hPXRx Mice in the DSS-Induced Colitis Model.** Unlike rifampicin, rifaximin is a nonsystemic antibiotic not significantly absorbed into the circulation. The preventive role of rifaximin was assessed in the DSS-induced colitis model. No significant changes were observed in the colon after rifaximin treatment alone in hPXRx, WT, and Pxr-null mice compared with the control groups (Supplemen-
tal Fig. 3). In contrast, DSS treatment increased crypt and epithelial cell damage, infiltration of granulocytes and mononuclear immune cells, and tissue edema in hPXR mice. It is noteworthy that rifaximin pretreatment significantly attenuated DSS-induced colon damage in hPXR mice compared with DSS treatment alone (Fig. 2A), as noted by an observed decrease in epithelial loss and mild inflammation. Rifaximin did not protect DSS-induced colitis in wild-type and Pxr-null mice as assessed by colon histology. In addition, several clinical parameters of colitis were statistically improved with rifaximin pretreatment in hPXR mice, such as daily body weight loss, rectal bleeding, diarrhea, and colon length compared with DSS administration alone (Fig. 2B). Meanwhile, there was no protection in WT and Pxr-null mice treated with DSS and rifaximin compared with DSS alone as revealed by clinical indexes of colitis, including body weight, rectal bleeding scores and diarrhea scores, and colon length (Supplemental Fig. 4). These results suggest that the beneficial effects of rifaximin may be attributed to its specific activation of human PXR, high bioavailability in the intestine, and decreased toxicity compared with rifampicin. To ensure that the protective role of rifaximin in acute colitis is not via inducing hepatic SCD-1, Western blot analysis of hepatic microsomal SCD-1 was performed. Rifaximin alone had no effect on SCD-1 expression in hPXR, wild-type, or Pxr-null mice, whereas 2.5% DSS administration clearly repressed SCD-1 expression in hPXR, wild-type, or Pxr-null mice compared with control (Supplemental Fig. 5A). Moreover, serum metabolomics analysis demonstrated that rifaximin-treated hPXR, wild-type, and Pxr-null mice have similar profiles of score plots and loading plots as determined by principal components analysis (Supplemental Fig. 5B), suggesting that rifaximin may be acting locally in the colon and not systemically to decrease the severity of DSS-induced colitis.

**Rifaximin Protects TNBS-Induced IBD in hPXR mice.** To confirm results from the DSS-induced colitis model, an additional acute colitis model was used. The TNBS-induced IBD model was driven by an initial IL-12 and TNFα response of lymphoid cells (Strober et al., 2009). The TNBS-induced IBD model was initiated in hPXR mice by intrarectal administration of TNBS after rifaximin pretreatment. TNBS triggered weight loss, bloody diarrhea, rectal prolapse, and large bowel wall thickening. Rifaximin significantly improved colon histology (Fig. 3A) and body weight (Fig. 3B) and maintained colon length (Fig. 3C) in hPXR mice, whereas no protection was observed in WT and Pxr-null mice (data not shown). Thus, with a second mouse IBD model, these data further indicate that rifaximin protects against IBD in hPXR mice.

**Rifaximin Exerts Therapeutic Efficacy in the DSS-Induced Colitis Model in hPXR Mice.** In addition to a preventive role of rifaximin, a therapeutic study was performed in DSS-induced hPXR and Pxr-null mice. Acute colitis was induced by 2.5% DSS treatment for 5 days. Rifaximin (1 mg/kg/day) was administrated to hPXR and Pxr-null mice after the onset of IBD. Survival rates were markedly increased in hPXR mice (P = 0.011) compared with Pxr-null mice (P = 0.730) administered rifaximin after the withdrawal of DSS (Fig. 4A). Clear signs of recovery from colitis were noted in hPXR mice administered rifaximin (Fig. 4B), such as increased colon length (P = 0.002) and increased body weight compared with controls. A significant resolution of colon damage was observed in hPXR mice treated with DSS alone compared with colon sections from rifaximin-treated
hPXR mice (Fig. 4C). No significant protection was afforded in 
Pxr-null mice treated with rifaximin with respect to sur-

daive (Fig. 4D) or improvement of clinical signs (Fig. 4E)

compared with hPXR mice.

Rifaximin Activates hPXR Target Genes in the Co-
lon. Human PXR expression in colons was measured by qPCR
(Fig. 5A) and Western blot (Fig. 5B). Mouse b-actin served as an
internal control for qPCR mRNA analysis, and HNF4b served as
an internal control for Western blot analysis. Human PXR was
expressed in the colons of hPXR mice; however, the expression was
significantly less than that observed in the liver. Previous work
demonstrated that rifaximin is a human PXR agonist in the small
intestine (Ma et al., 2007b). To assess the potential function of
rifaximin as a PXR ligand in the colon, qPCR analysis of mouse
Cyp3a11, Cyp3a13, Gsta1, and Mdr1a genes in colon was inves-
tigated. The expression of Cyp3a11, Cyp3a13, Gsta1, and Mdr1a
mRNAs was increased in hPXR mice after rifaximin treatment
compared with the control group (Fig. 6), whereas there was no
induction of Cyp3a11, Cyp3a13, Gsta1, and Mdr1a mRNAs in
wild-type and Pxr-null mice after rifaximin treatment. Together,
these data demonstrate that rifaximin is a human PXR agonist in
the colon.

Activation of hPXR by Rifaximin Inhibits NF-kB Tar-
gene expression. NF-kB is the central transcription
factor in the regulation of proinflammatory cytokines and
chemokines. Previous studies demonstrated that activation of
PXR can attenuate NF-kB signaling (Xie and Tian, 2006).
To elucidate the potential function of NF-kB on protection of
DSS-induced colitis by rifaximin, qPCR analysis of several
NF-kB-regulated proinflammatory cytokines and chemokines
in colon were evaluated. The results demonstrated that
mRNAs encoding the inducible nitric oxide synthase (iNOS),
chemokine motif receptor 2 (CCR2), TNFb, interferon-b
(IFNb), intercellular adhesion molecule 1 (ICAM-1), mono-
cyte chemotactant protein 1 (MCP-1), and interleukin
(IL)-10, IL-6, and IL-1b mRNA were induced in colonic tissue
after 5-day DSS treatment of hPXR mice compared with the
control group (Fig. 7). The increases in inflammatory medi-
ators (iNOS, CCR2, TNFb, IFNb, ICAM-1, MCP-1, IL-10, IL-
6, and IL-1b) after DSS treatment were significantly de-
creased in hPXR mice pretreated with rifaximin. These data
suggest that rifaximin-activated human PXR may protect
DSS-induced colitis through inhibiting the NF-kB-mediated
proinflammatory response.

Direct Inhibition of NF-kB by Rifaximin. To determine the
direct role of human PXR in the repression of NF-kB
signaling by rifaximin, luciferase activities of an NF-kB
reporter construct were measured. In HT-29 and Caco-2 cells,
TNFα (10 ng/ml) induced NF-κB activation, which was significantly attenuated after cotransfection with human PXR and RXR constructs. These data revealed that human PXR could inhibit TNFα-induced inflammation. The addition of 1 and 100 μM rifaximin potentiated the repression of NF-κB activation after TNFα treatment (Fig. 8A). The significant inhibition of NF-κB activity after rifaximin treatment in the absence of human PXR transfection might be caused by basal expression of human PXR in HT-29 and Caco-2 cells as assessed by Western blot analysis (data not shown). Moreover, a dose response of rifaximin on NF-κB inhibition was evaluated in HT-29 cells. These results showed that the optimal inhibition dosage of rifaximin was between 0.01 and 1 μM (Fig. 8B). Consistent with the in vivo results, the in vitro luciferase assay indicated that rifaximin protected hPXR mice from acute colitis through NF-κB inhibition via gut-specific human PXR activation.

**Rifaximin Inhibits NF-κB Target Genes through a hPXR-Mediated Mechanism.** NF-κB target genes iNOS, CCR2, TNFα, IFNα, ICAM-1, MCP-1, IL-10, IL-6, and...
deviations.

n/H9251 H9252

HNF4α cally recognizes human PXR but not mouse PXR or other liver proteins. The monoclonal antibody against human PXR (H5017) specifically recognizes human PXR but not mouse PXR or other liver proteins. HNF4α was used as a loading control.

Fig. 5. Basal expression of human PXR in the colon of hPXR mice. A, qPCR analysis of human PXR expression in colon and liver of hPXR mice and in colon of WT and Pxr-null mice. Mouse β-actin mRNA served as an internal control. B, Western blot analysis of human PXR expression in colon and liver of hPXR mice and in colon of WT and Pxr-null mice. Pooled nuclear protein (30 μg) was loaded for each sample (n = 3 per group). The monoclonal antibody against human PXR (H5017) specifically recognizes human PXR but not mouse PXR or other liver proteins. HNF4α was used as a loading control.

Fig. 6. mRNA analysis of hPXR target genes in colon tissue. Expression of mRNAs encoding Cyp3a11, Cyp3a13, Gsta1, and Mdr1a were determined by qPCR from colon epithelial cells isolated from hPXR, WT, and Pxr-null mice treated with control (Cont) and rifaximin (Rifax). Data were normalized to β-actin. Each bar represents the mean ± standard deviations. n ≥ 6, *, p < 0.05.

IL-1β mRNAs were measured in wild-type and Pxr-null mice after DSS treatment. A significant induction of all proinflammatory mediators was observed in wild-type and Pxr-null mice compared with the control group. However, rifaximin did not decrease inflammatory mediators after DSS treatment (Fig. 9), but actually increased TNFα and CCR2 mRNAs in Pxr-null mice and TNFα, IL-1β, IL-10, and MCP-1 in wild-type mice compared with DSS and rifaximin cotreatment to DSS treated alone, which might result from a differential inflammation response of the above factors in Pxr-null and wild-type mice. These data demonstrate that rifaximin affords protection mainly through a gut-specific hPXR-dependent mechanism and not through its antibiotic properties.

Discussion

The etiology of IBD is still not completely understood; however, it is widely accepted that the development of IBD is associated with the interplay of genetic, bacterial, and environmental factors and dysregulation of the intestinal immune system (Kaser et al., 2010). Among the IBD models, chemically induced animal models are commonly used, particularly with DSS colitis and hapten-induced colitis, using either DNBS/TNBS or oxazolone. These two models produce robust and immediate inflammation in the colon and have high reproducibility (Hoffmann et al., 2002). In the present study, acute colitis was induced by DSS and TNBS administration. Despite the acute nature of these models, they clearly show the impact of human PXR and rifaximin and suggest its clinical utility.

The treatment of IBD is highly individualized; however, in most cases the first-line therapy uses immune-suppressing drugs, which act systemically. Rifaximin, an efficient gut-specific antibiotic because of its zwitterionic property, which inhibits absorption into the systemic circulation (Marchi et al., 1985), has been shown to be beneficial in specific cases of IBD (Gionchetti et al., 2006). Using a mouse model that accurately expresses human PXR, the present study clearly demonstrates that rifaximin protects through a PXR-dependent mechanism and not as a general antibiotic in both DSS- and TNBS-induced IBD. In the therapeutic study, no mice in the DSS group survived, whereas rifaximin administered after treatment with DSS resolved the colitis symptoms and led to 60% survival. The resolution of inflammation in acute models of colitis is not completely understood; however, these data provide evidence that rifaximin and hPXR might be worthwhile for the treatment of human IBD.

It is noteworthy that a well known human PXR agonist, rifampicin, which is a systemic antibiotic used to treat tuberculosis, demonstrated an increase in hepatotoxicity when administered before treatment with DSS and provided no significant protection of hPXR mice in the DSS-induced colitis model. Rifampicin treatment in the colitis model actually slightly increased the severity of symptoms such as body weight loss and diarrhea score. This was quite unexpected because pregnenolone-16α-carbonitrile, which also induces PXR target genes in liver and colon, protects against DSS-induced IBD (Shah et al., 2007). The reason for the differences between the protective effect of rifaximin and rifampicin in experimental IBD is not understood. Because both drugs can induce PXR target genes in the intestine, the difference is not caused by the increase in the availability of rifaximin in the gut compared with rifampicin. Further experimentation is required to determine the reason for the lack of effect of rifampicin on DSS-induced colitis in the hPXR mouse. It was reported that rifampicin led to several pseudomembranous colitis cases (Akbar et al., 2003; Mazokopakis et al., 2008; Chen et al., 2009) and has been associated with adverse effects in patients with chronic inflammatory diseases (Yuhas et al., 2009). It is noteworthy that rifampicin administered with DSS markedly lowered the already suppressed SCD-1 protein and the serum levels of unsaturated fatty acids compared with DSS alone. The current results in
the mouse IBD model suggest that these clinical observations could be the result of the lowering of SCD-1 expression and associated anti-inflammatory unsaturated fatty acids (Chen et al., 2008).

Previous studies have shown that PXR is remarkably divergent across mammalian species with the ligand binding domains sharing 70 to 80% identity compared with the 90% typically exhibited by other nuclear receptors (Zhou et al., 2008).
2009). Because mouse PXR was reported to ameliorate DSS-induced IBD by NF-κB inhibition, it was critical to investigate the mechanism by which human PXR was protective in an acute colitis model. In vivo data demonstrated several NF-κB target genes to be significantly attenuated in hPXR mice after rifaximin pretreatment and DSS treatment compared with DSS treatment alone, and the attenuation of NF-κB signaling by rifaximin was not observed in wild-type and Pxr-null mice. Consistent with the in vivo data, in vitro evaluation of rifaximin on NF-κB inhibition suggested a direct role for human PXR in the inhibition of NF-κB signaling. Analysis of NF-κB response in colon-derived cancer cells using the NF-κB response element demonstrated a significant induction with TNFα treatment, which was inhibited after overexpression of human PXR, and the inhibition was further potentiated in the presence of rifaximin. It was assumed that transrepression of NF-κB target genes might be mediated by the SUMOylation of PXR (Zhou et al., 2006b), and NF-κB disrupted binding of the PXR–RXR complex to its binding motif (Gu et al., 2006).

In summary, the hPXR mice were used to demonstrate the beneficial effects of rifaximin in acute models of colitis through the activation of human PXR, and a critical role for PXR in IBD. In addition, these data provide a mechanistic basis for novel therapies targeting human PXR in the treatment of inflammatory diseases of the gut.

Acknowledgments
We thank John Buckley for assistance with the animal studies and Salix Pharmaceuticals, Inc., for supplying rifaximin.

References


Address correspondence to: Frank J. Gonzalez, Laboratory of Metabolism, Center for Cancer Research, National Cancer Institute, Building 37, Room 3106, Bethesda, MD 20892. E-mail: fjgonz@helix.nih.gov
Therapeutic Role of Rifaximin in Inflammatory Bowel Disease: Clinical Implication of Human Pregnan X Receptor Activation

Jie Cheng, Yatrik M. Shah, Xiaochao Ma, Xiaoyan Pang, Toshiya Tanaka, Tatsuhiko Kodama, Kristopher W. Krausz, Frank J. Gonzalez

SUPPLEMENTAL DATA
Supplementary Figure 1. Hepatic SCD-1 expression and serum metabolomics through LC-MS-based metabolomics in rifampicin-treated hPXR mice. (A) SCD-1 expression was evaluated by qPCR and Western blot. Liver tissues were isolated from hPXR mice treated with vehicle (Cont), DSS treatment alone (DSS), RIF (3 mg/kg/day) treatment alone (RIF3), RIF (10 mg/kg/day) treatment alone (RIF10), DSS treatment following RIF (3 mg/kg/day) pre-treatment (DSS/RIF3) and DSS treatment following RIF (10 mg/kg/day) pre-treatment (DSS/RIF10). qPCR analysis was used to measure SCD-1 mRNA, normalized to β-actin. For measurements of SCD-1 protein levels, liver microsomes were prepared by differential centrifugation and pooled microsomal samples (n>=3 livers in each group, 15 μg total loading proteins) were used for western blot analysis. GAPDH was used as a loading control. PosC: positive control from untreated liver microsomes (5 μg) of C57BL/6 mice. (C) Scores plot of PCA models (upper panel) from Cont, DSS, RIF3, RIF10, DSS/RIF3 and DSS/RIF10. A two-component PCA model was constructed to characterize the relationship among the treatment groups, Cont ▲, RIF3 ●, RIF10 □, DSS ■, DSS/RIF3 ▼, DSS/RIF10 ♦. The t [1] and t [2] values represent the scores of each sample in principal component 1 and 2, respectively. Fitness ($R^2$) and prediction power ($Q^2$) of this PCA model were 0.677 and 0.429. Loadings plot of chemical ions (below figure) were shown with pc [1] and pc [2] values, which represented the contributing weights of each ion to principal components 1 and 2 of the PCA model, respectively.
Supplementary Figure 2. Messenger RNA analysis of hPXR target genes and pro-inflammatory mediators in liver and colon. RNA was isolated from liver and colon of hPXR mice treated with Cont, DSS, RIF3, RIF10, DSS/RIF3 and DSS/RIF10. Expression of mRNAs encoding Cyp3a11, Cyp3a13, Gsta1 Mdr1a, iNOS, CCR2, TNFα, IFNα, ICAM-1, MCP-1, IL-10, IL-6 and IL-1β were determined by qPCR. Data was normalized to β-actin. Each bar represents the mean ± standard deviations, n>6 (* for p < 0.05, ** for p < 0.01, *** for p < 0.001).
Supplementary Figure 3. Rifaximin does not alter colon histology. Representative hematoxylin and eosin (H&E)-stained colon sections of hPXR, wild-type (WT) and Pxr-null treated with control (Cont) and rifaximin (Rifax). Magnification: 200 X.
Supplementary Figure 5. Colitis assessment of wild-type (WT) and Pxr-null male mice. WT and Pxr-null mice were treated with control (Cont), rifaximin (Rifax), DSS and DSS/rifaximin (DSS/Rifax) and body weight, diarrhea, rectal bleeding and colon length were assessed in (A) WT and (B) Pxr-null mice. Each bar represents the mean ± standard deviations, n>=6 (* for p < 0.05, *** for p < 0.001).
Supplementary Figure 4. SCD-1 expression and serum metabolomics in rifaximin-treated hPXR, wild-type (WT) and Pxr-null mice. (A) SCD-1 levels in liver microsomes were evaluated by western blot analysis. Liver tissues were isolated from hPXR, WT and Pxr-null mice treated with control (Cont), DSS, rifaximin (Rifax) and DSS/rifaximin (DSS/Rifax). Liver microsomes were prepared by differential centrifugation. Pooled microsomal samples (n=6 livers in each group, 15 μg total loading proteins) were used for Western Blot analysis. GAPDH was used as a loading control. PosC: positive control from untreated liver microsome (5 μg) of C57BL/6 mice. (B-D) Scores plot of PCA models (upper figures) from hPXR, WT and Pxr-null mice. A two-component PCA model was constructed to characterize the relationship among the treatment groups, Cont ▲, Rifax ○, DSS ■ and DSS/Rifax ◆. The t [1] and t [2] values represent the scores of each sample in principal component 1 and 2, respectively. Fitness ($R^2$) and prediction power ($Q^2$) of this PCA model were 0.631 and 0.157 for hPXR, 0.660 and 0.392 for WT, 0.760 and 0.253 for Pxr-null, respectively. Loadings plot of chemical ions (below figures) from serum samples of hPXR, WT and Pxr-null mice were shown with pc [1] and pc [2] values, which represented the contributing weights of each ion to principal components 1 and 2 of the PCA model, respectively. Major ions contributing to the separation of DSS/Rifax from other groups were labeled as C18:0-LPC, C18:1-LPC, C18:2-LPC, and C22:6-LPC, respectively.