Slc10a2-null mice uncover colon cancer-promoting actions of endogenous fecal bile acids

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Abstract

Although epidemiological evidence in humans and bile acid feeding studies in rodents implicate bile acids as tumor promoters, the role of endogenous bile acids in colon carcinogenesis remains unclear. In this study, we exploited mice deficient in the ileal apical sodium-dependent bile acid transporter (ASBT, encoded by SLC10A2) in whom fecal bile acid excretion is augmented more than 10-fold. Wild-type and Asbt-deficient (Slc10a2⁻/⁻) male mice were treated with azoxymethane (AOM) alone to examine the development of aberrant crypt foci, the earliest histological marker of colon neoplasia and a combination of AOM and dextran sulfate sodium to induce colon tumor formation. Asbt-deficient mice exhibited a 54% increase in aberrant crypt foci, and 70 and 59% increases in colon tumor number and size, respectively. Compared to littermate controls, Asbt-deficient mice had a striking, 2-fold increase in the number of colon adenocarcinomas. Consistent with previous studies demonstrating a role for muscarinic and epidermal growth factor receptor signaling in bile acid-induced colon neoplasia, increasing bile acid malabsorption was associated with M3 muscarinic and epidermal growth factor receptor expression, and activation of extracellular signal-related kinase, a key post-receptor signaling molecule.

Introduction

Primary bile acids are synthesized in the liver from cholesterol, N-acyl amidated to taurine or glycine, secreted into bile and, in most mammals, stored in the gallbladder before meal-stimulated release into the proximal small intestine where they facilitate lipid digestion. Although small quantities of bile acids are reabsorbed passively throughout the gut, ~95% of intestinal bile acid uptake is mediated in the distal small intestine by the ileal apical sodium-dependent bile acid transporter (ASBT, encoded by SLC10A2) (1). Bile acids absorbed from the gut return to the liver via the portal venous circulation and are re-excreted into bile and the small bowel. This enterohepatic circulation minimizes bile acid loss in stool. ASBT-deficiency markedly impairs bile acid absorption and fecal bile acid excretion is greatly augmented in Asbt-deficient (Slc10a2⁻/⁻) compared to wild-type (WT) male mice (2).

Although chronic or intermittent diarrhea is the most overt clinical manifestation of bile acid malabsorption, the long-term consequences of persistently exposing colon epithelial cells to increased bile acid levels are uncertain. One concern is that by direct or indirect actions on colon epithelial cells fecal bile acids promote colon neoplasia. Whereas colon cancer arises from progressive mucosal dysplasia associated with somatic gene mutations, environmental factors can accelerate and augment this process. Diets rich in meat and fat increase both fecal bile acid levels and colon cancer risk (3).
Surprisingly little is known about ASBT expression and activity in colon cancer. Wang et al. (4) reported a greater than 2-fold increase in the risk of colon adenomas in persons with SLC10A2 gene polymorphisms, but a case–control study found no increase in colon cancer risk with common SLC10A2 gene variants (5). These studies were limited by small sample sizes, failure to control for diet, drugs, cholecystectomy and other factors that modulate intestinal bile acid levels and composition, and did not correlate SLC10A2 gene polymorphisms with either tumor size or progression. As growth factors, bile acids are more likely to act as tumor promoters than initiators (6–8).

In rodent models of chemical carcinogen-induced colon neoplasia, instillation of exogenous bile acids into the gut promotes epithelial dysplasia and tumor formation (3). Even without a genetic predisposition to colon neoplasia or exposure to a carcinogen, mice fed a human secondary bile acid, deoxycholic acid, for 8–10 months developed colon cancer (8). Collectively, these observations provide evidence that excess levels of endogenous bile acids can promote colon neoplasia. In contrast, the effects of endogenous bile acids on the development of chemical carcinogen-induced colon neoplasia in rodents have only been examined in surgical models where large portions of the small bowel were resected or bypassed to increase bile acid flux into the colon. An important limitation of those studies is that bowel resection disrupts normal intestinal function and results in malabsorption of many luminal constituents in addition to bile acids. Indeed, studies using that approach found increased chemical carcinogen-induced colon neoplasia in rats following resection of either the proximal small intestine or the distal small intestine (9,10). To overcome the limitations of surgical approaches we used Asbt-deficient mice as an in vivo model of bile acid malabsorption and to assess the cancer-promoting potential of chronically exposing colon epithelial cells to elevated levels of endogenous bile acids. In Asbt-deficient mice, the lack of bile acid uptake in the distal small intestine, which we recently documented using novel in vivo imaging (11,12), results in more than 10-fold increased fecal bile acids (2). Unlike surgical models, fecal weight is not increased in Asbt-deficient mice despite a significant decrease in colon transit time (2).

### Materials and methods

#### Animals

We used male mice with targeted disruption of the Asbt gene (Slc10a2−/− mice) and wild-type (WT) male mice, all backcrossed more than 10 generations on a C57BL/6 genetic background (2). For all experiments, mice were housed under identical conditions in a pathogen-free environment with a 12:12-h light/dark cycle and free access to water and standard mouse chow (the Teklad Global 18% Protein Extruded Rodent Diet contains as percentage of calories: 24% protein, 18% fat and 58% fiber). All animal studies were conducted in accordance with the Guide for the Care and Use of Laboratory Animals prepared by the U.S. National Academy of Sciences. Studies conducted at Wake Forest University School of Medicine were approved by that school’s Institutional Animal Care and Use Committee. Animal studies conducted in Baltimore were approved by both the Institutional Animal Care and Use Committee at the University of Maryland School of Medicine and the Research and Development Committee at the VA Maryland Health Care System.

### Quantitative real-time polymerase chain reaction

To examine the expression of intestinal transcripts, total RNA from tumors and adjacent normal colon was extracted from tissue stored in RNAlater. Five micrograms of RNA were used for first-strand complementary DNA synthesis using the Superscript III First Strand Synthesis System for RT-PCR (Invitrogen). Quantitative real-time polymerase chain reaction was performed with SYBR Green PCR Master Mix (Applied Biosystems) and specific primers at a final concentration of 0.5 μM in sample volumes of 20 μL. The primer sequences shown in Table 1 were designed to span introns (National Center for Biotechnology Information nucleotide database SIM-4 gene alignment program and online software; http://www.genetools.com/index.html). qPCR was performed using the 7900HT Fast System and SYBR Green Master Mix (ABI) following the manufacturer's protocol. PCR conditions were 5 min at 95°C, followed by 37 cycles of 95°C for 15 s, 60°C for 20 s and 72°C for 40 s and a final cycle at 95°C for 15 s. Data were analyzed using ABI instrument software (SDS 2.1). Expression of GAPDH was used as an endogenous control for normalization. All samples were run in triplicate and the Ct (2−ΔΔCt) method was used to compare samples when assessing the relative expression of transcripts.

### Immunohistochemical staining

To identify and quantify protein expression, formalin-fixed paraffin-embedded tissue sections were immunostained with specific antibodies against M3 muscarinic receptors (M3R) (Alomone Labs, Jerusalem, Israel), extracellular signal-related kinase (ERK) and phospho-ERK (Cell Signaling Institute, Danvers, MA). Immunohistochemical staining was performed as described previously (13).

### Abbreviations

<table>
<thead>
<tr>
<th>Abbreviation</th>
<th>Definition</th>
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<tr>
<td>ACF</td>
<td>aberrant crypt foci</td>
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<tr>
<td>AOM</td>
<td>azoxymethane</td>
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<tr>
<td>ASBT</td>
<td>apical sodium-dependent bile acid transporter</td>
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<tr>
<td>DSS</td>
<td>dextran sulfate sodium</td>
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<tr>
<td>ERK</td>
<td>extracellular signal-related kinase</td>
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<tr>
<td>FXR</td>
<td>farnesoid X receptor</td>
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<td>M3R</td>
<td>M3 muscarinic receptors</td>
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<td>MMP7</td>
<td>matrix metalloproteinase-7</td>
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<td>WT</td>
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Abbreviations adapted from (14).
Ki67 staining

To examine cell proliferation, Ki67 staining was performed as described by Holt et al. (14). Only the distal half of the colon and complete crypts were analyzed by investigators masked to genotype and treatment. To measure changes in the distribution of Ki67 staining, crypts were divided equally into five zones, starting at the base of the crypt (zone 1) and extending to the luminal surface (zone 5). The ‘labeling index’ was calculated as the percentage of cells in a crypt zone that stained for Ki67 (14); a minimum of 1000 colonocytes was evaluated per mouse colon.

Statistics

Data are presented as average value ± standard error of the mean. The unpaired Student’s t test (assuming unequal variance) was used for comparing continuous variables between two independent groups. A logarithm transformation (log(tumor volume + 1)) was applied to tumor volume so that the transformed variable approximately followed a normal distribution. Then, because multiple tumors could be present in one mouse, a linear transformation (Log(tumor volume + 1)) was applied to tumor volume so that the transformed variable approximately followed a normal distribution. Differences in body weight between male WT and Asbt-deficient mice treated with AOM (7.5 mg/kg by intraperitoneal injection once each week for 4 weeks) plus 2.5% DSS (added to the drinking water for 5 days) were compared using Student’s t test (assuming unequal variance) followed by post hoc tests with Tukey–Kramer’s post hoc tests.

Results

ACF provide the earliest histological evidence of colon neoplasia, sometimes appearing in murine models within a few weeks of carcinogen treatment (13). To investigate the effects of Asbt deficiency on colon neoplasia, we examined ACF formation in WT and Slc10a2−/− mice treated with AOM, a commonly used experimental carcinogen (6). Starting at age of 4 weeks, 11 WT and 10 Slc10a2−/− mice received 7.5 mg/kg AOM by intraperitoneal injection once each week for 10 weeks. Mice were killed 20 weeks after the first AOM injection and their colons were harvested, stained with methyl blue and examined for ACF. ACF were identified microscopically in methylene blue-stained harvested colon tissue by readily identifiable features including enlarged crypts, a thickened layer of methylene blue-stained epithelial cells, slit-shaped luminal openings, increased pericytchal space and elevation above the focal plane (Figure 1A, inset) (13). We found a statistically significant, 60% increase in the number of ACF in colons from AOM-treated Slc10a2−/− mice compared to those from WT mice (P < 0.05; Figure 1A).

Even though they may develop ACF, some mouse strains, like C57BL/6, resist progression to colon tumor formation following treatment with AOM alone (3), a limitation circumvented by adding DSS to their drinking water. We treated mice using the study design illustrated in Figure 1B: 20 WT and 15 Slc10a2−/− male littermates were treated with AOM (7.5 mg/kg by intraperitoneal injection once each week for 4 weeks) plus 2.5% DSS (added to the drinking water for 5 days). Four WT and three Slc10a2−/− mice treated with vehicle alone served as additional controls. As shown in Figure 1B, Asbt-deficient mice weighed approximately 10% less than WT littermates; this significant difference (P < 0.05) persisted throughout the study period, including a transient drop in body weight immediately after DSS treatment; three Slc10a2−/− mice died within a week of DSS treatment. The small differences in body weight between male Slc10a2−/− and WT mice, reported previously (2), did not impact colon length (data not shown).

Although no colon tumors were detected in vehicle-treated mice, most WT and Asbt-deficient animals developed tumors (Figure 1C). As shown in Figure 1C and D, Asbt deficiency was associated with a 70% increase in the mean number of colon tumors per mouse (P = 0.02) and 59% increase in mean tumor size (P = 0.04). Asbt deficiency also significantly increased the multiplicity of colon tumors; nine WT mice versus only one Asbt-deficient mouse had fewer than two tumors (P = 0.049, Fisher’s two-tailed exact test; Figure 1C).

Increased cell proliferation is a hallmark of neoplastic transformation and bile acids stimulate colonocyte proliferation in vitro (15). To examine the actions of increased fecal bile acids on colonocyte proliferation in vivo, we measured Ki67 immunostaining (Figure 2A). As shown in Figure 2B, Slc10a2−/− differentiated ACF Ki67 immunostaining depending upon the crypt zone examined; in colons from Slc10a2−/− compared to those from WT mice there were significantly fewer Ki67-stained cells at the crypt base (zone 1); the reverse was observed at the crypt apex (zone 5) (both P < 0.05). The shift of proliferating cells to the outer portions of the crypt in Slc10a2−/− mice likely represents an early manifestation of dysplasia. We measured colonocyte apoptosis using caspase-3 immunostaining; as observed previously (6) caspase-3-stained cells were an order of magnitude fewer than Ki67-stained cells and there were no significant differences in either normal epithelial or tumor cell apoptosis.
when comparing WT and Asbt-deficient mice (data not shown). These findings support the concept that fecal bile acids act primarily as growth factors that promote cell proliferation.

Mouse colon tumors represented both adenomas and adenocarcinomas; two Asbt-deficient mice, but no WT mice, developed sessile-serrated adenomas (Figure 2C), a biomarker for activation of epidermal growth factor receptor signaling (16). Notably, as shown in Figure 2D, the increased tumor burden in Asbt-deficient mice resulted from more adenocarcinomas, not adenomas. Whereas the overall number of adenomas was unchanged, Asbt-deficient mice had a nearly 2-fold increase in the number of adenocarcinomas (P < 0.002).

In the colon, two bile acid transporters expressed on the colonocyte basolateral membrane mediate export of passively absorbed bile acids into the enterohepatic circulation; hetero-meric Organic Solute Transporter (OSTβ-OSTβ) and multidrug resistance-associated protein-3 (MRP3) (17). Bile acid induced changes in OSTβ-OSTβ expression are regulated in part by the bile acid nuclear receptor farnesoid X receptor (FXR) (17). To explore the consequences of increased fecal bile acids, we compared the expression of genes for FXR and these bile acid transporters in tissue from Asbt-deficient to that from WT mice.

In humans and murine colon neoplasia models, expression of the nuclear bile acid receptor FXR is reduced in colon carcinoma compared to adjacent normal mucosa (18). Thus, it was not surprising to find that in adenocarcinomas from both WT and Asbt-deficient mice Fxr expression was significantly reduced compared to normal colon (Figure 3A). Similar changes were observed for the genes encoding Mrp3 (Abcc3), Ostα and Ostβ (Figure 3A), most likely reflecting regulation of bile acid transporter expression by FXR.

To gain additional insight into the mechanisms underlying the promotion of colon neoplasia in Asbt-deficient mice, we examined the relative expression of genes commonly associated with colon cancer progression. As shown in Figure 3B, expression levels of the genes for cyclooxygenase 2 (Ptgs2), cMyc and cyclinD1 (CyclD1) in both WT and Asbt-deficient mice were significantly increased in adenocarcinomas. However, there was no significant difference in the relative expression of these genes in normal colon and tumors from Asbt-deficient mice when compared with those from WT mice (solid black and red bars in Figure 3B). This suggests that altered expression of Ptgs2, cMyc, Ctnnb1 and CyclD1 is most probably a consequence of neoplastic transformation independent of Asbt deficiency.

Consistent with its important role in colon cancer progression (7), expression of the gene for matrix metalloproteinase-7 (Mmp7) was strikingly induced in murine colon tumors; 40- to 400-fold in tumors versus adjacent normal colon from WT mice and Asbt-deficient mice (P < 0.001; Figure 3B). Moreover, in normal tissue from Asbt-deficient compared to WT mice, Mmp7 expression was increased 8-fold (P < 0.01; Figure 3B), suggesting...
that induction of Mmp7 expression in normal colon may be relevant to bile acid-induced promotion of colon neoplasia.

Muscarinic receptors and ligands play important roles in colon neoplasia (6,7); M3R are over-expressed in colon cancer and their activation stimulates colon cancer cell proliferation. In animal models of genetic and sporadic colon cancer, M3R knockout or inactivation attenuates murine colon epithelial cell proliferation and neoplasia (6,19). We used immunohistochemistry to examine M3R expression in normal colon epithelium and carcinomas from WT and Asbt-deficient mice. As illustrated in Figure 3C and D, left panel, in both WT and Asbt-deficient mice we observed significantly increased M3R staining in colon cancers compared to normal tissue and a more modest, but statistically significant, increase in M3R expression in tumors from Asbt-null compared to WT mice (Figure 3D, right panel; \(P < 0.05\)). In tumors from Asbt-deficient animals, we detected nearly 2-fold increased expression of Chrm3, the gene encoding M3R (Figure 4A), a finding indicative of M3R activation (20). Chrm3 over-expression in Asbt-deficient mice was of similar magnitude to that observed when AOM-treated WT mice were fed bethanachol, a muscarinic receptor agonist (21); bethanachol treatment had also stimulated 30 and 52% increases in tumor number and size (21), of similar magnitude to what we observed in Asbt-deficient mice (Figure 1C and D). Moreover, like Asbt deficiency (Figure 2D), bethanachol treatment increased the numbers of colon adenocarcinomas, not adenomas (21).

In seeking evidence of post-M3R signal transduction, we detected up-regulation of mRNA for epidermal growth factor receptor in tumors from Asbt-deficient mice (Figure 4A), and in both WT and Asbt-deficient mice, immunostaining for epidermal growth factor receptor protein was increased in tumors compared to normal tissue (immunohistochemical staining scores: 2.88 ± 0.18 versus 2.04 ± 0.17 for WT mice; 3.05 ± 0.09 versus 1.90 ± 0.16 for Asbt-deficient mice; both \(P < 0.005\)) (Figure 4B). Whereas normal tissue and tumors from WT and Asbt-deficient animals had similar levels of total ERK, in both genotypes levels of phospho-ERK were significantly increased in tumors
compared to normal tissue (Figure 4B and C). Notably, in tumors from Asbt-deficient mice we detected a modest but statistically significant increase in ERK phosphorylation (P < 0.05; Figure 4D), supporting increased ERK activation.

Discussion

In the current study, we used mice deficient in the predominant intestinal bile acid transporter, ASBT, to show that increased spillage of endogenous bile acids into the feces augments the number of ACF and tumors in the colon, and increases tumor size. Notably, in Asbt-deficient mice, the number of adenocarcinomas, not adenomas, was increased. This observation of advanced neoplasia in Asbt-deficient mice supports the concept that increased fecal bile acids stimulate tumor growth, not initiation. This concept also agrees with the natural history of the Asbt-deficient mouse; over the period 2002–2013 we maintained 522 Asbt-deficient mice (295 on a C57BL/6 genetic background) to our knowledge, this is the first study using a non-surgical model to suggest that endogenous bile acids can promote colon neoplasia in vivo. Although bile acids such as deoxycholate were shown to induce formation of reactive oxygen and nitrogen species and oxidative DNA damage, and can stimulate muscarinic receptor signaling in colon-derived cell lines, we recognize that evidence in the present study is insufficient to prove that bile acids were directly responsible for the increased neoplasia in

Previously, using human colon cancer cells, we showed that bile acids activate M3Rs and stimulate cross-talk with epidermal growth factor receptors by a mechanism involving activation of MMP7 (22); muscarinic receptor activation robustly induces MMP7 mRNA expression (7,21). In fact, in Asbt-deficient mice we detected robust overexpression of MMP7 mRNA in both normal colon tissue as well as colon tumors. Since crosstalk between M3 muscarinic and epidermal growth factor receptors is a prominent feature of proliferative signaling in human colon cancer cells (15), our finding of activated ERK signaling in Asbt-deficient mouse colon provides additional support for a mechanistic interaction between M3 muscarinic and epidermal growth factor receptors in mediating the actions of increased fecal bile acids (15). Similar outcomes in mice that ingested bethanachol, a muscarinic receptor agonist, and Asbt-deficient mice with increased spillage of bile acids into the colon lends additional support to this proposed mechanism. Collectively, these observations support bile acid-induced activation of muscarinic receptor signaling as a mechanism whereby increased fecal bile acids induce Mmp7 expression and promote colon neoplasia in Asbt-deficient mice.

To our knowledge, this is the first study using a non-surgical model to suggest that endogenous bile acids can promote colon neoplasia in vivo. Although bile acids such as deoxycholate were shown to induce formation of reactive oxygen and nitrogen species and oxidative DNA damage, and can stimulate muscarinic receptor signaling in colon-derived cell lines, we recognize that evidence in the present study is insufficient to prove that bile acids were directly responsible for the increased neoplasia in

Figure 3. Increased Mmp7 gene expression, and muscarinic and EGF receptor expression, and activation of post-receptor signaling in tumors from Slc10a2−/− mice. (A) qRT-PCR of colon epithelial cell bile acid nuclear receptor and transporters. Expression of genes for Fxr, Mrp3 (Abcc3) and Ostβ was significantly reduced in adenocarcinomas from WT and Slc10a2−/− mice compared to adjacent normal colon (*P < 0.05, **P < 0.01, ***P < 0.001; n = 9–11). (B) qRT-PCR of markers for colon cancer progression. Expression of Psy2, cMyc, CycD1 and Mmp7 was increased in adenocarcinomas from WT and Slc10a2−/− mice compared to adjacent normal colon (*P < 0.05, **P < 0.01; n = 9–11). Mmp7 expression was significantly increased in normal colon from Slc10a2−/− mice compared to normal colon from WT mice (**P < 0.01; ***P < 0.001). (C) Representative images of M3R staining in normal colon tissue and adenocarcinomas from WT and Slc10a2−/− mice. (D) M3R staining is increased in tumors compared to normal colon epithelium from both WT and Asbt-deficient mice (left panel). NI, normal tissue. Bars represent mean ± SEM for staining intensity score measured in 0.5 increments by a senior pathologist on a scale from 0, no staining to 3, maximal staining (**P < 0.01; n = 10–13). M3R staining was significantly increased in tumors from Asbt-deficient (KO) mice compared to those from WT mice (right panel). Bars represent mean ± SEM for staining intensity expressed in arbitrary units (**P < 0.05; n = 10–13).
vivo. Genetic ablation of Asbt is much more selective than intestinal resection, but bile acid malabsorption is likely to also affect the absorption of other small molecules. As such, the concentration of other endogenous and dietary metabolites may be altered in the soluble phase of the colonic contents. It is also noteworthy that changes in bile acid spectrum and concentration can impact the composition of the gut microbiota and vice versa. Since there is emerging evidence that changes in gut microbial metabolism can affect the development of colon cancer in genetically susceptible models or after treatment with tumor initiators such as AOM/DSS, it is also possible that bile acid malabsorption-associated changes in the gut microbiota may play a role in development of colon neoplasia (23–25).

Despite these limitations, our findings have potentially important clinical implications. Recent studies suggest that idiopathic bile acid malabsorption is not uncommon in the general population and may be a major cause of chronic diarrhea and diarrhea-predominant irritable bowel syndrome. Bile acid malabsorption is also common in inflammatory bowel disease involving the terminal ileum and bariatric surgical procedures such as Roux-en-Y gastric bypass which may increase bile acid flux through the colon. Bile acid diarrhea is a side effect of chemotherapeutic agents such as lenalidomide, and may be an underappreciated effect of several classes of FDA-approved drugs which inhibit bile acid transport by ASBT in vitro (26). Finally, drugs that specifically inhibit ASBT are being developed to treat hypercholesterolemia and constipation; chronic use of such agents may unintentionally promote colon neoplasia.

**Supplementary Material**

Supplementary Figure 1 can be found at [http://carin.oxfordjournals.org/](http://carin.oxfordjournals.org/)

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Conflict of Interest Statement: PAD is listed as the inventor on two patents owned by Wake Forest University related to screening technology for ASBT inhibitors. PAD has previously served as a consultant for Lumena Pharmaceutical and GlaxoSmithKline. The other authors report no conflicts.

References