Products of the Colonic Microbiota Mediate the Effects of Diet on Colon Cancer Risk\textsuperscript{1,2}

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Abstract

It is estimated that most colon cancers can be attributed to dietary causes. We have hypothesized that diet influences the health of the colonic mucosa through interaction with the microbiota and that it is the milieu interior that regulates mucosal proliferation and therefore cancer risk. To validate this further, we compared colonic contents from healthy 50- to 65-y-old people from populations with high and low risk, specifically low risk Native Africans (cancer incidence <1:100,000; \( n = 17 \)), high risk African Americans (risk 65:100,000; \( n = 17 \)), and Caucasian Americans (risk 50:100,000; \( n = 18 \)). Americans typically consume a high-animal protein and -fat diet, whereas Africans consume a staple diet of maize meal, rich in resistant starch and low in animal products. Following overnight fasting, rapid colonic evacuation was performed with 2 L polyethylene glycol. Total colonic evacuants were analyzed for SCFA, vitamins, nitrogen, and minerals. Total SCFA and butyrate were significantly higher in Native Africans than in both American groups. Colonic folate and biotin content, measured by \textit{Lactobacillus rhamnoses} and \textit{Lactobacillus plantarum} ATCC 8014 bioassay, respectively, exceeded normal daily dietary intakes. Compared with Africans, calcium and iron contents were significantly higher in Caucasian Americans and zinc content was significantly higher in African Americans, but nitrogen content did not differ among the 3 groups. In conclusion, the results support our hypothesis that the microbiota mediates the effect diet has on colon cancer risk by their generation of butyrate, folate, and biotin, molecules known to play a key role in the regulation of epithelial proliferation.


Introduction

Population studies have shown wide variations in the risk of colon cancer. Differences in inherited susceptibility characteristics and related gene-environment interactions are some of the possible explanations. Diet-gene interactions have been the most extensively studied and are among the most widely accepted explanations for the variation. Colon cancer is the second leading cause of cancer death in Americans, and African Americans are at the highest risk of developing and dying from colon cancer. However, our recent studies have highlighted the paradox that Africans living in Africa rarely get this disease (1,2). The observation that colon cancer is more common in developed than in developing nations suggests that the westernized way of life may be responsible for the increased risk of the disease (3). The aspect that has attracted the most attention is diet, as experimental studies have provided robust evidence for the ability of nutrients to modify cancer risk. Some dietary factors, such as fresh fruit and vegetables, fiber, antioxidants, and calcium and vitamin D suppress risk, whereas others such as red meat and saturated animal fat increase risk (3,4). Based on our own investigations, we have concluded that high intakes of red meat may be responsible for the increased risk in African Americans, whereas the low incidence in Africans may be a consequence of their high intakes of maize (corn) meal and low consumption of meat and animal fat (1). A novel aspect of our investigations was the identification of major differences in colonic bacterial metabolism. In African Americans, undigested carbohydrate was chiefly metabolized by hydrogen-producing bacteria, whereas in Africans it was mainly fermented by methane-producing bacteria (1). Our conclusion that the colonic microbiota was different was backed up by our detection of higher rates of colonization with secondary bile salt-producing bacteria in African Americans and \textit{Lactobacilli} species in Africans. It is established that a diet high in animal fat stimulates the growth of secondary bile salt-producing bacteria and further studies have shown that secondary bile salts are cytotoxic and carcinogenic (5). A diet rich in red meat also stimulates the growth of sulfate-reducing bacteria, which produce hydrogen sulfide, which experimentally is known to be genotoxic (6). In contrast, \textit{Lactobacilli} promote mucosal health (7,8), as do butyrate-producing \textit{Firmicutes}. There is extensive experimental evidence that the SCFA butyrate is essential for colonic mucosal health. It is the preferred energy source for colonocytes and it

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suppresses epithelial proliferation, which is a recognized biomarker for cancer risk. Finally, the microbiota synthesize essential vitamins and recent studies have shown that transport mechanisms for their absorption exist in the colon. Two of these, folate and biotin, are intimately involved with epigenetic regulation of epithelial proliferation.

Amalgamating these findings, we developed the hypothesis that the differences in cancer risk may be attributed to differences in the colonic milieu, which represents the microbial interface between the external environment (namely the diet) and the colonic mucosa (9). Here, we examine differences in the constituents of the colonic milieu that have been shown experimentally to influence neoplastic change from high and low risk populations.

Methods

Study populations
As colon cancer predominantly occurs in middle life and after, colonic content samples from African American and Native African men and women aged 50–65 y were examined. As a western control, Caucasian Americans of a similar age range were included. The intention was to recruit by advertisement in public fora 20 participants of either sex from each population without history of gastrointestinal disease or surgery or colon polyps or cancer and without a history of antibiotic use during the previous 8 wk. This age group was selected, because the risk of cancer increases appreciably after age 50 y and screening colonoscopy is recommended as the standard of care. Americans were recruited predominantly from the Pittsburgh area of Pennsylvania and the Africans from semiurban Pretoria (Garankuwa, Medical University of South Africa), central highlands) and from the rural regions of Limpopo Province, South Africa. The protocol was reviewed and approved by the University of Pittsburgh Institutional Review Board and General Clinical Research Center and by the Medical University of South Africa’s Medical Ethics and Safety Committee. The protocol for collection of samples in South Africa was the same as that used at the University of Pittsburgh.

Experimental design
Participants consumed their normal diets until 2000 h and then fasted until the study was complete. At 0800 a blood sample (10 mL) was taken from fasting participants and then they were given 2 L of colonoscopy preparation fluid (polyethylene glycol 3550, GoLytely) to drink rapidly from fasting participants and then they were given 2 L of colonoscopy preparation fluid (polyethylene glycol 3550, GoLytely) to drink rapidly over 15–30 min. All stool passed during the following 3 h was collected in preweighed cans and kept on ice. Visual inspection showed that by this time, the residual effluent had little fecal contamination. The can was then reweighed to calculate the total weight of the stool passed. The container was sealed and emulsified by rigorous shaking for at least 30 min. The fluid was then carefully inspected to ensure there was no solid material remaining and 30-mL aliquots were extracted and frozen at −20°C to await transportation to the laboratory for analysis. Transportation from Africa to Pittsburgh was performed by packaging frozen samples on dry ice and by rapid airfreight transfer (<3 d). Visual inspection showed that samples were still frozen on arrival at the investigator’s laboratory in Pittsburgh.

The results of the dietary analysis, colonoscopic findings, epithelial proliferation rates, and microbiota differences have been previously published (1).

Sample preparation

Colonic fluid for SCFA and vitamins. Duplicates of ~5 g were emulsified for 2 min and then centrifuged at 3800 × g for 5 min at 5°C. The supernatant was decanted, split in 4, and frozen at −20°C until analysis.

Colonic minerals. Five grams of stool was dried thoroughly by centrifugation under vacuum (CentriVap Concentrator, Labconco) for analysis.

Other vitamins. Twelve evacuant samples from each group were pooled, diluted with saline, centrifuged at 3800 × g for 5 min, and refiltered at 200 nm for a pilot investigation into the colonic fluid content of vitamin B-12, thiamine, and vitamin C.

Blood tests. A 5-mL sample was collected in an EDTA tube for determining the complete blood count and another 5 mL collected without anticoagulant for measuring plasma folate concentrations.

Sample analysis

SCFA. Concentrations were measured by GC (Agilent Technologies 6890N Network GC System with FID fitted with EC-1000 15-mL 0.53 mm × 1.2 mm ECONO-CAP capillary column) based on the method described by Scheppach et al. (10). A mixed SCFA standard solution was made up from reagents of the highest purity (>99%) (Sigma). 2,2-Dimethylbutyric acid at a concentration of 1 mmol/L was used as an internal standard. SCFA values were computed using peak area ratio of the analyte to the internal standard based on the standard curve. There was a good linear correlation between the peak area ratio and the corresponding standard SCFA with r values > 0.99 for all SCFA. The inter-day and intra-day CV ranged from 2.4 to 3.9%.

Colonic folate. Folate concentrations in fecal fluid were measured by 2 methods in a subset of 10 Africans and 10 African Americans by chemiluminescence folate affinity assay using a competitive binding assay on an automated Centaur Instrument (Siemens) in our hospital laboratory (i.e. the method used for blood folate analysis) and then by specific microbial bioassay (11) in Dr. O’Connor’s laboratory to first assess the contribution of other fecal contaminants that register false positive values. Despite a poor correlation between colonic folate measured by the affinity assay and the bioassay, the results were of a similar order of magnitude [mean ± SE 769 ± 153 μg vs. 412 ± 57 μg, respectively (folate 2.266 ng/ml = 1 mmol/L)]. Some samples gave very similar values, whereas others were 4X higher by the affinity assay. Consequently, the bioassay method (duplicate variation <1%) was used in the final analysis shown in the results section.

Microbial bioassay. Aliquots of fecal fluid fraction supernatants were diluted with 1% sodium ascorbate in potassium phosphate buffer (pH 7) to protect labile folates, heated (100°C for 5 min) to remove folate from binders, and after cooling, folates were converted to their microbiologically assayable form using rat serum folate conjugase as previously described (12). The folate concentrations in the resultant supernatants were then determined by microbiological assay described by Molloy and Scott (11) using the test organism Lactobacillus rhamnosus (ATCC 7469, American Type Tissue Culture Collection).

Colonic biotin. An improved agar plate method of biotin bioassay using Lactobacillus plantarum ATCC 8014 and bromocresol purple was established to determine biotin levels in body fluids, foods, and feces (13). Samples were treated with 4.5 mol/L H2SO4 to liberate free biotin, autoclaved for 1 h, and neutralized by 4.5 mol/L NaOH, then 10 mL was added to wells in each plate. The intra-assay CV were 3.2 (n = 20) and 1.3% (n = 23), respectively.

Thiamine. Total thiamine contents were determined as sum of thiamine and thiamine diphosphate. Trichloroacetic acid (5%) was added to the sample and supernatant of the mixture was used for measurement. Thiamine and thiamine diphosphate were determined by the HPLC-post labeled fluorescence method as previously described (14).

Ascorbic acid. Reduced and oxidized ascorbic acid and 2,3-diketogulonic acid were chemically converted to ascorbic acid derivative and the compound was determined by the HPLC method (15).

Vitamin B-12. The assay was run on the Bayer Centaur 1556 (Advia Centaur, Siemens) with calibration lot CC12 and reagent lot 191. The assay is a competitive solid phase immune assay with chemiluminescence detection (16). The assay was repeatable to within 5% in the sample range (reagent lot 191; calibration lot CC13).

Colonic minerals. Following sample digestion with nitric acid and treatment with hydrogen peroxide and hydrochloric acid (17), calcium,
iron, zinc, and sulfur were measured in the resulting digestion solution by sequential inductively coupled plasma optical emission spectrometry (17) (Perkin-Elmer Optima 2000 DV ICP-OES, PerkinElmer Life And Analytical Sciences) utilizing VHGI multi-element aqueous standard SM35A-500, trace-metal grade ammonium sulfate (Sigma-Aldrich), and Standard Reference Material 1577b (National Institute of Standards and Technology). Nitrogen was determined by the Kjeldahl method (17).

**RBC folates.** EDTA-preserved whole blood samples (50 μL) were used for analysis of folate concentrations following hemolysis with ascorbic acid (1 mL) in the hospital laboratory using a similar method (chemiluminescence folate-affinity assay) described above for colonic samples. Measurements were converted to RBC folate concentrations after correction for hematocrit using the formula:

\[
\text{RBC folate (mg/L)} = \frac{\text{folate result for hemolysate, mg/L} \times 21}{100 \text{ hematocrit}}
\]

where 21 is the dilution factor (conversion factor: 2.266 ng/ml = 1 nmol/L).

**Statistical analysis**
As the data derived from the stool analysis were normally distributed, significant differences among the 3 population groups were first assessed by 1-way ANOVA. If there was an overall difference among the colonic contents of the 3 different population groups, then comparisons were made using Fischer’s post hoc protected least significant difference test and the adjusted significance value for multiple testing. Regression analysis was used to assess the association between dietary intakes and nutrient colonic contents. Values in the text are means ± SEM. Differences were considered significant at \( P < 0.05 \).

**Results**

**Dietary intakes.** Dietary intake results have been previously published (1) (Table 1). Native Africans consumed a relatively impoverished diet, significantly lower in protein, vitamins, and minerals than in both other groups.

**Colonic evacuants.** Total colonic evacuant volumes were slightly, but significantly, higher in African Americans compared with Africans, but not compared with Caucasian Americans (Table 2). Consequently, the differences in colonic mineral content between African Americans and native Africans would be smaller if fecal concentrations alone were examined, whereas the differences in SCFA would be higher.

**Colonic minerals and nitrogen.** The calcium and iron contents were significantly higher in Caucasian Americans, whereas that of zinc was significantly higher in African Americans than in Native Africans (Table 2). Unfortunately, assessment of normal basal fecal levels of sulfur could not be made, because the colonic evacuant used, GoLytely, contained 1280 mg/L elemental sulfur. Colonic nitrogen contents were similar despite the considerably higher dietary protein intakes in Americans.

**Colonic SCFA.** The quantities of all 3 major SCFA, namely acetate, propionate, and butyrate, were all higher in native Africans than in African Americans and Caucasian Americans, with the exception of acetate, which tended to be lower in Caucasian Americans than in native Africans (\( P = 0.07 \)) (Fig. 1). There was no significant difference between Colonic SCFA contents in African Americans and Caucasian Americans did not differ. Because colonic volume was lower in Africans, the concentrations of SCFA in colonic effluent were even more elevated (\( P < 0.001 \)) in Africans compared with both American groups (data not shown). Overall, there were no significant correlations between colonic butyrate and total dietary carbohydrate and fiber intake.

**Colonic folate.** The colonic folate content was not lower in Africans despite the lower dietary intake (201 ± 23 μg/d) than African Americans (481 ± 47 μg/d) (\( P < 0.001 \)) and Caucasian Americans (526 ± 50 μg/d) (\( P < 0.0001 \)) and lower RBC folate concentrations than in native Africans (Table 2). The low RBC concentrations in Africans were not, however, associated with functional evidence of deficiency as indicated by anemia or macrocytosis. There was a weak overall association between dietary carbohydrate and dietary fiber intake and colonic folate content (\( r^2 = 0.18, P = 0.02 \); and \( r^2 = 0.17, P = 0.04 \)). The association within the 2 American groups for folate and dietary carbohydrate was stronger (\( r^2 = 0.26, P = 0.01 \)).

**Colonic biotin.** Large quantities of biotin were also recovered from colonic effluents in all 3 groups (Table 2). The colonic content in Caucasian Americans (183 ± 68 μg) was similar to the U.S. recommended dietary intake (RDA) of 150–300 μg/d, but the quantity measured in native Africans exceeded considerably the measured dietary intake levels, i.e. 91 ± 24 μg compared with only 19 ± 4 μg/d (1). Colonic biotin concentrations correlated positively with colonic folate concentrations (\( r = 0.56; P < 0.0001 \)).

**TABLE 2**

<table>
<thead>
<tr>
<th></th>
<th>Native Africans</th>
<th>African Americans</th>
<th>Caucasian Americans</th>
</tr>
</thead>
<tbody>
<tr>
<td>( n )</td>
<td>17</td>
<td>17</td>
<td>18</td>
</tr>
<tr>
<td>Colon contents</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Weight, g</td>
<td>1302 ± 108</td>
<td>1796 ± 181*</td>
<td>1694 ± 226</td>
</tr>
<tr>
<td>Calcium, mg</td>
<td>683 ± 119</td>
<td>1084 ± 143</td>
<td>1944 ± 260**</td>
</tr>
<tr>
<td>Nitrogen, mg</td>
<td>3131 ± 497</td>
<td>2459 ± 390</td>
<td>3054 ± 443</td>
</tr>
<tr>
<td>Iron, mg</td>
<td>289.4 ± 4.0</td>
<td>30.1 ± 6.2</td>
<td>39.2 ± 4.9*</td>
</tr>
<tr>
<td>Zinc, mg</td>
<td>6.1 ± 1.4</td>
<td>20.4 ± 5.2*</td>
<td>13.4 ± 5.2</td>
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<tr>
<td>Folate, μg</td>
<td>632 ± 95</td>
<td>699 ± 131</td>
<td>860 ± 129</td>
</tr>
<tr>
<td>Biotin, μg</td>
<td>91 ± 24</td>
<td>65 ± 21</td>
<td>183 ± 58</td>
</tr>
<tr>
<td>Blood</td>
<td></td>
<td></td>
<td></td>
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<tr>
<td>Hb, g/d</td>
<td>131 ± 10</td>
<td>138 ± 3</td>
<td>141 ± 3</td>
</tr>
<tr>
<td>MCV, fl</td>
<td>85 ± 4</td>
<td>90 ± 2</td>
<td>93 ± 2*</td>
</tr>
<tr>
<td>RBC folate, ng/mL</td>
<td>181.3 ± 20.0</td>
<td>328.2 ± 34.2**</td>
<td>341.4 ± 41.4**</td>
</tr>
</tbody>
</table>

1 Values are means ± SEM. Asterisks indicate different from Native Africans: * \( P < 0.05 \), ** \( P < 0.005 \).

2 Reproduced from (1) with permission.
Insignificant, whereas vitamin B-12 (AA 2.5 mg intake levels. RDA 1 mg) show that the colonic content of butyrate, and all of the chief antineoplastic properties (19–22). Butyrate does not come from the diet but is produced by specific colonic bacteria, predominately members of the Clostridia clusters XIVa and IV of Firmicutes (23), from food residues such as dietary fiber or resistant starch. Our results show that the colonic content of butyrate, and all of the chief SCFA, were significantly higher in native Africans, the group that has the lowest colon cancer risk, i.e. < 1:100,000, compared with 65:100,000 in African Americans and 50:100,000 in Caucasian Americans (1). Our results also illustrate the potentially vital role the colonic microbiota may play in vitamin synthesis in undernourished communities, such as the rural Africans included in this study who had higher quantities of folate and biotin in their colons than in their usual diet. Despite the fact that participants had fasted 12 h, we cannot conclude that the total quantities of vitamins recovered from the colon were microbiota-derived, because dietary vitamin absorption by the small intestine is never 100%. The ability of the microbiota to produce folate. For example, Pompei et al. (33) found that among rats with 3 of these bacteria increased plasma and liver folate concentrations and that the effect was amplified by the further dietary supplementation with fructo-oligosaccharides, which functions similarly to resistant starch and fiber as a prebiotic (32).

There remains concern, however, that body folate status was marginal in Africans, because RBC levels, an accepted measure of body stores, were low, but not low enough to produce anemia and macrocytosis. We speculate that the rich topical source of folate in the colon may suppress neoplastic change as there is experimental evidence that localized folate deficiency plays a key role in premalignant changes in the epithelium (31). The importance of topical, or local, folate as opposed to systemic blood levels for mucosal health might also explain the other side of the coin: why rural Africans have one of the highest risks of squamous cell carcinoma of the esophagus in the world (34), whereas the cancer is rare in Americans.

Equally impressive was the colonic content of biotin, which is also a product of bacterial fermentation and therefore likely induced by dietary residues. Little attention has been paid to biotin’s potential role in carcinogenesis. The vitamin was...
thought to be principally involved in carbohydrate metabolism as a coenzyme for carboxylases involved in fatty acid synthesis, amino acid catabolism, and gluconeogenesis (13). However, recent investigations have suggested that biotin may exert similar epigenetic effects as folate, because biotinylatation, like methylation, of histones has been shown to regulate cell proliferation through its influence on DNA transcription, replication, and repair (35). Another similarity in function between folate and biotin is that marginal tissue deficiencies of both are associated with birth defects (36). Could it be that combined biotin and folate topical sufficiency in the colon suppresses colon cancer risk?

In summary, our study supports our hypothesis that colon cancer risk is determined by interactions between the diet and the colonic microbiota and that cancer risk can therefore be modified by dietary and microbial manipulation. The failure of short-term dietary interventions to prevent adenomatous polyp recurrence and cancer may be explained by the multiple protective mechanisms of the body to regulate DNA turnover and the fact that it takes a lifetime to develop environmentally induced cancers. Consequently, short-term dietary intervention studies need to first prove that biomarkers of cancer risk, e.g. epithelial proliferation, can be changed as proof of concept before longitudinal interventional programs are introduced to reduce cancer risk in high risk populations, such as African Americans.

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Literature Cited