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Morphological adaptation with preserved proliferation/transporter content in the colon of patients with short bowel syndrome

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Joly F, Mayeur C, Messing B, Lavergne-Slove A, Cazals-Hatem D, Noordine ML, Cherbuy C, Duée PH, Thomas M. Morphological adaptation with preserved proliferation/transporter content in the colon of patients with short bowel syndrome. *Am J Physiol Gastrointest Liver Physiol* 297: G116–G123, 2009. First published April 23, 2009; doi:10.1152/ajpgi.90657.2008.—In short bowel syndrome (SBS), although a remaining colon improves patient outcome, there is no direct evidence of a mucosal colonic adaptation in humans. This prospective study evaluates morphology, proliferation status, and transporter expression level in the epithelium of the remaining colon of adult patients compared with controls. The targeted transporters were Na⁺/H⁺ exchangers (NHE2 and 3) and oligopeptide transporter (PepT1). Twelve adult patients with a jejuno-colonic anastomosis were studied at least 2 yr after the last surgery and compared with 11 healthy controls. The depth of crypts and number of epithelial cells per crypt were quantified. The proliferating and apoptotic cell contents were evaluated by revealing Ki67, PCNA, and caspase-3. NHE2, NHE3, PepT1 mRNAs, and PepT1 protein were quantified by quantitative RT-PCR and Western blot, respectively. In patients with SBS compared with controls, 1) hyperphagia and severe malabsorption were documented, 2) crypt depth and number of cells per crypt were 35% and 22% higher, respectively ($P < 0.005$), whereas the proliferation and apoptotic levels per crypt were unchanged, and 3) NHE2 mRNA was unmodified; NHE3 mRNA was downregulated near the anastomosis and unmodified distally, and PepT1 mRNA and protein were unmodified. We concluded that, in hyperphagic patients with SBS with severe malabsorption, adaptive colonic changes include an increased absorptive surface with an unchanged proliferative/apoptotic ratio and well-preserved absorptive NHE2, NHE3, and PepT1 transporters. This is the first study showing a controlled nonpharmacological hyperplasia in the colon of patients with SBS.

nutrition; colonic hyperplasia; proliferation markers; Ki67; PCNA; caspase-3; Na⁺/H⁺ exchanger 3; Na⁺/H⁺ exchanger 2; PepT1; hyperphagia

THE NORMAL HUMAN SMALL INTESTINE length is generally considered to be between 3 and 8 m from the duodenojejunal flexure to the ileocecal valve. In adults, the massive resection of the small bowel leaving less than 200 cm of small bowel defines short bowel syndrome (SBS), and a small bowel length of <100 cm is highly predictive of permanent intestinal failure (8, 31). SBS includes clinical signs characterized by diarrhea, dehydration, malabsorption of macronutrients, weight loss, malabsorption of vitamins and trace elements, and malnutrition (2, 37). In patients with SBS, the degree of malabsorption depends on the area and site of the resection as well as any

persistence of damage within the remnant bowel. Patients with SBS type I had a jejunostomy, whereas patients with SBS type II had an anastomosis between jejunum and colon.

After massive reduction of the small bowel in animal models, the remnant intestine shows functional changes and significant morphological changes, i.e., hyperplasia, to compensate for absorptive functions in both the small bowel and the colon (18–20, 27, 44, 48). These changes are called adaptive and are mainly dependent on enteral neuro-hormonal stimuli (53). This adaptive phase occurs between the postoperative and sequela phases both in animals and humans. In humans, an adaptive process has never been convincingly demonstrated in the remnant small or large intestine (3, 39, 41). At the colonic level, absorptive changes in humans are significant through hyperfermentation of malabsorbed carbohydrates attributable to changes in the bacterial microflora (7). Indeed, clinical experience indicates that patients with short bowel with a jejuno-colonic anastomosis (SBS type II) show improved absorption capacity with time, whereas patients with end-terminal jejunostomy without colon (type I) do not (12, 31), and the presence of remaining colon is associated with a lower dependence of a supply with parenteral nutrition (PN) (31, 35). Although the presence of the colon is an important determining factor in the clinical SBS outcome, with a better outcome than type I SBS (36, 51), direct proof of mucosal colonic adaptive changes have not been demonstrated except for one study showing an increase in H⁺-coupled oligopeptide transporter (PepT1) in the remaining colon (55).

The aim of this prospective study was to evaluate the nutritional status and the adaptive changes of the remaining colon of SBS adult patients, at least two years after resection, compared with healthy controls.

MATERIALS AND METHODS

A prospective case-control study was performed. The Human Investigations Committee of Saint Louis Hospital-Paris approved the study (no. 031048).

Patients

Inclusion criteria of patients with SBS were 1) SBS type II with a postduodenal remnant jejunum <150 cm, no remnant ileum and reestablished jejuno-colic continuity for more than 2 yr, 2) a nonrestricted oral diet, and 3) a normal colonoscopy of the remaining colon. All patients received gastric antisecretory drugs, but none had pancreatic enzyme treatment. Exclusion criteria were 1) upper gastrointestinal tract surgery (esophagus, stomach, duodenum, and/or pancreas), 2) a remnant ileum, 3) radiation enteritis and active Crohn's disease, 4) organ failure other than gastrointestinal, 5) evolutive neoplasia, 6) intestinal fistulae, 7) treatment by recombinant human

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growth hormone or glucagon like peptide-2 in the last 12 mo, 8) body mass index (BMI) >30, 9) age >80 or <18 yr old, 10) sepsis and/or antibiotics or aspirin in the 4 wk before the study, 11) treatments with steroids or immunosuppressive therapy in the 4 mo preceding the study, and 12) pregnancy. Reconstructive surgery of the remnant small bowel such as a reversed distal jejunal loop was not an exclusion criteria (31). The remaining colon in continuity, expressed in terms of percent of the usual length according to the method of Cummings et al. (13), was estimated on the basis of preoperative records. Routine biological data included serum albumin, transthyretin, and C reactive protein (CRP) blood levels.

Control Group

Participants were recruited among outpatients with an indication for colonoscopy for colon cancer screening or evaluation of abdominal pain. Control subjects were selected if they were clinically stable with no change in weight of more than 5% in the last 6 mo and had no previous history of gastrointestinal or systemic diseases. Other exclusion criteria included 1) BMI >30, 2) age >80 or <18 yr old, 3) sepsis and/or drugs in the 4 wk before the study, and 4) pregnancy. Inclusion criteria were a normal colonoscopy with normal histology sampling.

Absorption Study

Patients with SBS underwent a metabolic study to evaluate intestinal macronutrient absorption as described (32). Briefly, over a 6-day period, the first 3-day equilibrium period confirmed that patients were continuing their spontaneous intake of energy, carbohydrates, lipids, proteins, and fibers, and the absorption was measured the last 3 days. Unrestricted intake was measured by amount or weight and calculated with Bilnut software (Bourgerette P, Rolshansen M; BILNUT 4.0; SCDA Nutrisoft, Cerelles, France). Stools were collected daily and frozen at -20°C . Protein, lipid, and total energy were determined by nitrogen elemental analysis (N analyser Flash EA1112; Thermo Scientific, Waltham, MA) (26) and the method of Van de Kamer and bomb calorimetry (PARR 1351 bomb calorimeter; Parr Instrument, Moline, IL) (33), respectively. Quantification of carbohydrate-derived energy was calculated by subtracting the energy associated with the protein and lipid components from the total energy. The calorie-conversion factors used were 4.2, 9.35, and 5.65 kcal/g for carbohydrates, lipids, and proteins, respectively (47). The coefficient of net intestinal absorption represented the proportion of ingested energy not recovered in stool output.

Endoscopy and Mucosal Biopsy Specimens

To obtain tissue for histological analysis, patients and controls were submitted to routine bowel preparation with poly-ethylene-glycol the evening before colonoscopy with no change in dietary habits. Colonoscopy and colonic biopsies were performed in patients with SBS after the metabolic study. Three biopsies, allowed by the Human Investigations Committee, were obtained in the middle segments of remaining colon, i.e., proximal (or right colon), transverse, and distal colon (or left colon), respectively.

Analysis of Colonic-mucosal Morphology and Detection of Ki67 by Immunohistochemistry

All colonic biopsies were fixed in formalin for 24 h, embedded in paraffin, and analyzed on hematoxylin-eosin-safran-stained slides. Determination of epithelial cells and of depths of crypts were assessed by a senior gastrointestinal pathologist who was blind to the clinical findings (A. Lavergne-Slove) using a light microscope equipped with a calibrated ocular micrometer; each longitudinal hemicypt was divided into three equal-sized compartments from the muscularis mucosa to the surface. The quantifications required the full length of at least three well-oriented colonic crypts; this condition was obtained

in eight patients with SBS and seven controls. Immunohistochemistry was performed on deparaffinized formalin-fixed tissues sections (5 μm) using Ki67 antibody (clone Mib1; Dakocytomation, Glostrup, Denmark), and a histostainer automat (Nexes; Ventana, Tucson, AZ). The proliferative index of the mucosa was calculated by dividing the number of Ki67-labeled cells by the total number of cells and expressing the results as a percentage.

Detection of Caspase-3 by Immunohistochemistry

Caspase-3 immunohistochemistry was performed on deparaffinized formalin-fixed tissue sections (5 μm) with a rabbit monoclonal antibody against active caspase-3 (clone C92-605; BD Pharmingen, San Diego, CA) using a histostainer automat (Ventana). Caspase-3-positive cells were quantified as the higher number of positive colonic superficial cells observed between six crypts ($\times 200$ magnification), in eight patients with SBS and eight controls.

RNA Extraction and Real-time PCR Analysis

Frozen biopsy samples were homogenized using the TissueLyser (Retsch; Qiagen, Valencia, CA) for 5 min, 25 Hz in 1 ml Trizol reagent (TRIzol reagent; Invitrogen, Carlsbad, CA), and RNAs were isolated using the phenol-based extraction method (10). RNA concentration and purity were confirmed by absorbance measurement, using a Biophotometer (Eppendorf, Westbury, NY). In our experiment, A260:A280 ranged from 1.8 to 2.0, which is considered to be an acceptable indicator of RNA purity. An RNA integrity number (RIN) was also determined with the Agilent 2100 bioanalyzer (Agilent Technologies, Santa Clara, CA) and the RNA 6000 nano LabChip kit. The RINs did not exceed 5 in either biopsy. Because the RINs were close, samples were compared together as in Imbeaud et al. (23).

One microgram of the total RNA samples was subjected to a reverse transcription step in a 50- μl final volume using the high-capacity cDNA archive kit (Applied Biosystems, Foster City, CA). Single-stranded cDNA products were analyzed by real-time PCR using TaqMan Gene Expression Assays (Applied Biosystems). Single-stranded cDNA products were analyzed with the ABI PRISM 7000 Sequence Detection System (Applied Biosystems). Four genes were analyzed including: the eukaryotic 18S rRNA (Hs99999901_s1), the human NHE3 gene (SLC 9A3, Hs00188200_m1), the human NHE2 gene (SLC 9A2, Hs00268166_m1), and the human PEPT1 gene (SLC 15A1, Hs00192639_m1). After we tested several house-keeping genes, eukaryotic 18S rRNA was used as a reference gene because of its stability in the different colonic segments between patients and controls. In each case, triplicate cycle (Ct) values were obtained and averaged. The fold change in one target gene was normalized to the eukaryotic 18S rRNA.

Protein Extraction and Western Blot Analysis

Biopsies were disrupted with a tissue lyser in RIPA buffer, and total proteins were quantified using the Lowry method. Proteins were extracted from subconfluent HT-29 and Caco-2 line cells. Different amounts of proteins were loaded into a 10% SDS-polyacrylamide gel and then electrotransferred. PCNA and PepT1 were respectively revealed by a monoclonal antibody (Upstate Biotechnology, Lake Placid, NY) at a 1:1,000 dilution and by a polyclonal antibody (H-235; Santa Cruz Biotechnology, Santa Cruz, CA) at a 1:200 dilution. All membranes were stripped and rehybridized with an antibody against GAPDH (1:5,000) (Euromedex, Souffelweyersheim, France). An anti-mouse (1:12,000) or an anti-rabbit (1:5,000) antibody (Jackson ImmunoResearch Laboratories, Bar Harbor, ME) conjugated with peroxidase was used to detect immune complexes by chemiluminescence using the ECL+ detection system (Amersham Pharmacia Biotech, St. Giles, UK). Systematically, a protein ladder was loaded, and the weight of the band was checked. Autoradiography films were quantified using the Las1000 camera system (Fujifilm,

Table 1. Clinical characteristics of patients with SBS

Patient (sex)	Age When Studied, yr	Delay Since Continuity Reestablishment, yr	Diagnosis	Remnant Small Bowel Length, cm	Remnant Colon, %
P1 (F)	52	11	mesenteric infarction	0	70
P2 (M)	51	4	mesenteric infarction	10	50
P3 (F)	54	5	mesenteric infarction	10	80
P4 (F)	69	4	mesenteric infarction	15	80
P5 (F)	45	2	mesenteric infarction	20	80
P6 (M)	56	18	mesenteric infarction	30	40
P7 (M)	56	6	mesenteric infarction	35	80
P8 (M)	48	4	mesenteric infarction	35	100
P9 (M)	46	11	mesenteric infarction	40	50
P10 (M)	49	12	Crohn's disease	60	50
P11 (F)	71	9	volvulus	80	50
P12 (F)	60	19	mesenteric infarction	100	80
Means \pm SD	54 \pm 8	9 \pm 5		32 \pm 30	60 \pm 19

Patients were numbered (from 1 to 12) according to the remnant postduodenal jejunal length (cm). Delay was in years from the last surgery establishing an anastomosis between the jejunum and the remaining colon. The length of the remaining colon in continuity was expressed in percent of the usual length (13). SBS, short bowel syndrome; F, female; M, male.

Paris, France) and the Aida software (Raytest, Paris, France). The quantitative data (arbitrary unit) were calculated for each lane by dividing the signal of PepT1 by the signal of loading control.

Statistical Analysis

Statistical analysis was performed using the Statview software (version 5; SAS Institute, Cary, NC). The comparison between the control group and patients (means \pm SD) for age, BMI, oral intake, crypt depth, and number of cells per crypts was analyzed using a Student's *t*-test. $1/\Delta Ct$ values of RT-PCR were compared at each site (proximal, transverse, and distal) between control group and patients with a Mann-Whitney test. Level of significance was a *P* value <0.05 .

RESULTS

Characteristics of Patients with SBS and Controls

After obtaining written consent for the protocol, 12 patients with SBS and 14 controls were screened and eligible between January 2004 and September 2006.

Clinical characteristics of adult patients with SBS are listed in Table 1. All patients with SBS had a normal colonic mucosal histology. Etiology of SBS was arterial mesenteric infarction ($n = 10$), volvulus ($n = 1$), and Crohn's disease ($n = 1$) with no residual lesions of the remnant intestine. The only patient with Crohn's disease was in clinical remission without specific treatment for more than one year. The study was performed 9.0 ± 5.6 yr following reestablishment of bowel continuity (9.8 ± 5.7 yr following the small bowel resection). Patients with SBS had a total oral intake of $2,600 \pm 715$ kcal/day equivalent to 2.0 ± 0.46 times their resting energy expenditure (REE). Fecal wet weight was $1,780 \pm 540$ g/day (range 800–4,960 g/day) with a malabsorption of dietary protein,

energy, and fat (Table 2). Of 12 patients, nine were PN dependent. BMI was 18.5 ± 2.4 kg/m² with normal albumin 36 (34–47) g/l and transthyretin protein 210 (170–300) mg/l levels. The CRP was 1 (0.2–3.0) mg/l.

The age of controls (48.5 ± 15 yr) was not significantly different from that of patients with SBS. Mean oral intake was 1,457 (900–1,880) kcal/day in controls representing 1.0 ± 0.17 times REE, and BMI was 23 ± 2 kg/m² (Table 2). Oral intake and BMI of patients with SBS were respectively higher and lower ($P < 0.001$) than those of controls.

Morphological Data

Crypt depth was significantly ($P < 0.005$) increased ($+35 \pm 20\%$) in the colon of patients with SBS ($n = 8$, 464 ± 74 μ m) compared with controls ($n = 7$, 296 ± 45 μ m) (Fig. 1A). In accordance with this observation, the total number of epithelial crypt cells was also higher ($+22 \pm 16\%$) in patients with SBS ($P < 0.005$) than in control group (Fig. 1A). The proliferation rate of the remaining colon in patients with SBS was assessed with Ki67 as a marker of proliferative cells and caspase-3, as a surrogate of apoptosis. The ratio of Ki67-positive cells per crypt was not significantly different in patients with SBS ($n = 7$) and controls ($n = 7$) (Fig. 1B). A complementary analysis by Western blot indicated that PCNA, a protein involved in proliferation, was not modified after resection in the remnant colon (see Supplemental Fig. S1; supplemental material for this article is available at the *American Journal of Physiology Gastrointestinal and Liver Physiology* website). The apoptotic index was 9.8 ± 7 in patients with SBS ($n = 8$) and 9.2 ± 3 in controls ($n = 9$), indicating that the number of caspase-3-

Table 2. Intestinal absorption in patients with SBS and characteristics of controls

Patient (n = 12)	BMI, kg/m ²	Kcal/day	Oral Intake				Absorption				NP/week	REE, kcal/day	Intake/REE
			Protein, g	Fat, g	Glucide, g	Fiber, g	Total, %	Protein, %	Fat, %	Glucide, %			
	18.5 \pm 2	2,600 \pm 715	106 \pm 29	103 \pm 22	302 \pm 110	22 \pm 10	53 \pm 16	43 \pm 23	27 \pm 29	88 \pm 11	3.2 \pm 1.8	1,284 \pm 147	2.0 \pm 0.46
Control (n = 11)	23 \pm 2	1,457 \pm 75	69 \pm 18	48 \pm 15	182 \pm 32	15 \pm 7	Not Determined	Not Determined	Not Determined	Not Determined	No	1,471 \pm 232	1.0 \pm 0.17

Values are means \pm SD. Oral intake was assessed using Bilnut software. BMI, body mass index; REE, resting energy expenditure; intake/REE, ratio between oral intake (kcal/day) and REE (kcal/day); NP, number of parenteral nutrition infusions per week.

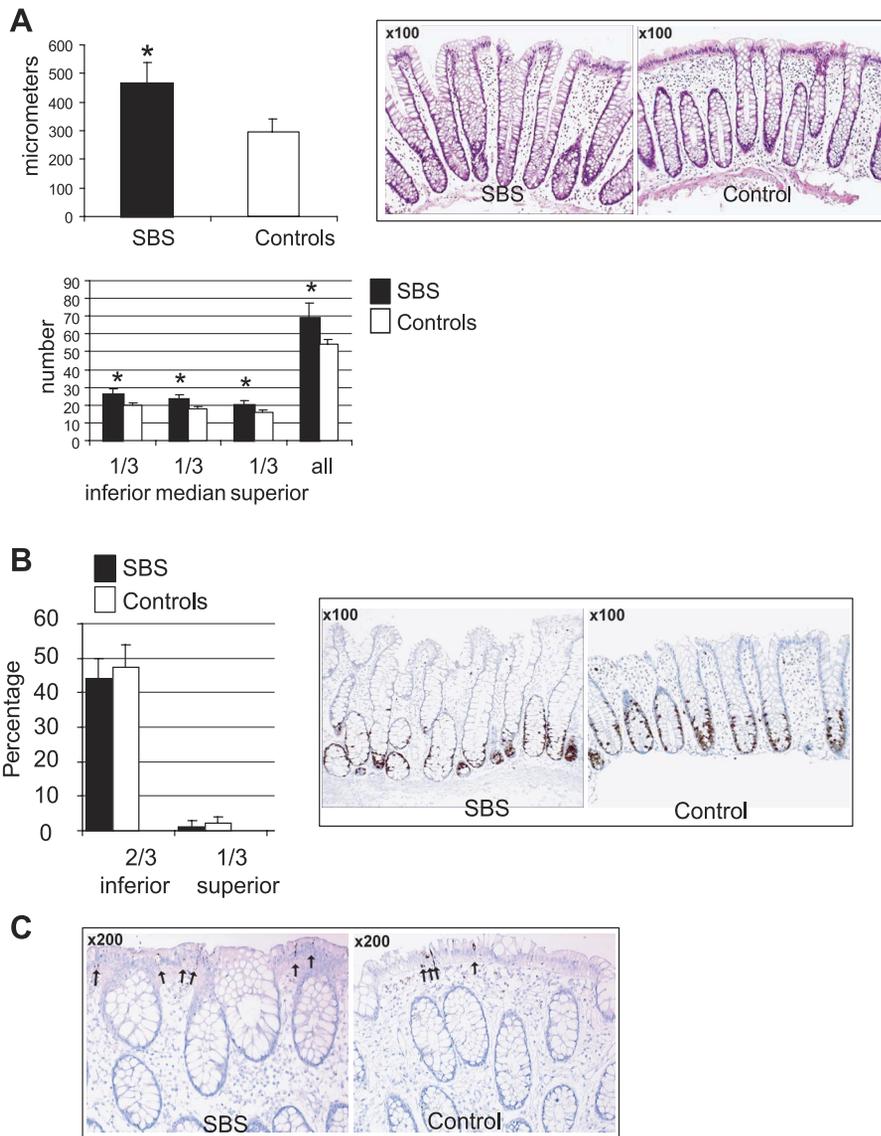


Fig. 1. Morphology in colon of patients with short bowel syndrome (SBS) and controls. *A*: morphology of colonic mucosa in patients with SBS compared with a control patient (hematoxylin and eosin). The number of cells per crypt was counted on longitudinal hemicypts divided into 3 equal-sized compartments (1/3 inferior, 1/3 median, 1/3 superior). * $P < 0.05$. *B*: percentage of Ki67-positive cells was expressed as a percentage in function of the total number of cells. *C*: caspase-3-positive cells were quantified as the positive colonic cells observed between 6 crypts (SBS and controls, $n = 8$).

positive cells was equivalent in patients with SBS and in controls (Fig. 1C).

NHE3, *NHE2*, and *PepT1* in the Colon of Patients with SBS and Controls

To evaluate the transport function of the remnant colon, we sought to analyze three genes, *NHE2*, *NHE3*, and *PepT1*. *NHE2* and *NHE3* play a major role in sodium and water absorption in the colon (4, 42, 54), whereas *PepT1* is involved in dipeptide uptake (mainly in the small intestine) (1). These genes were of particular interest because it has been proposed that *NHE2*, *NHE3*, and *PepT1* were overexpressed after a resection in animal models and in humans (34, 55).

Colonic expression hierarchy of NHE3, NHE2, and PepT1 transporters in controls. We determined *NHE3*, *NHE2*, and *PepT1* expression in the three colonic segments, i.e., proximal, transverse, and distal, of controls (Fig. 2). The hierarchy of colonic expression levels was $NHE3 > NHE2 > PepT1$. Throughout the anteroposterior gradient, *NHE3* was statistically more highly expressed in the proximal colon than in the

distal, whereas the opposite gradient of expression was observed for *PepT1*. *NHE2* was equally abundant from the proximal to the distal site.

mRNA of NHE3 and NHE2 exchangers in the colon of patients with SBS and controls. Because we observed that the transporters displayed a site-specific expression throughout the anteroposterior gradient, the level of mRNA was compared between patients with SBS and the control group for each proximal, transverse, and distal location (Fig. 3). The amount of *NHE3* mRNA was decreased at the proximal site ($P < 0.05$) in patients with SBS compared with controls, whereas it was not modified at the transverse or the distal sites (Fig. 3). The average of *NHE2* mRNA expression level was equivalent in patients with SBS and controls (Fig. 3).

PepT1 mRNA level and protein content in the colon of patients with SBS and controls. The average of *PepT1* mRNA expression level was equivalent in patients with SBS and controls (Fig. 3). However, there was a wide variability in the expression of *PepT1* mRNA in patients (data not shown). The variability of *PepT1* detected at the mRNA level in patients and

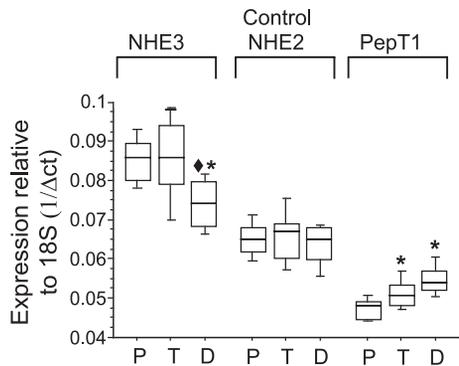


Fig. 2. Na^+/H^+ exchanger (NHE3 and NHE2) and PepT1 mRNA distribution in the control group. Eight controls (C1 to C8) were used for real-time RT-PCR analysis to detect NHE3, NHE2, and PepT1 mRNAs. The values were expressed as an average of $1/\Delta\text{Ct}$ [$1/(\text{Ct target gene}-\text{Ct18s})$]. T, transverse; P, proximal; D, distal site throughout the proximal-distal axis of the colon. ♦Statistically different from the transverse site ($P < 0.05$), *statistically different from the proximal site ($P < 0.05$).

to a lesser extent in controls, was not reflected at the protein level (Fig. 4A). There was no apparent increase in PepT1 protein in patients (P1, P2, P3, P5, and P6) compared with controls (C1, C2, C4, and C5). The PepT1 protein detection system was checked with cell lines and small bowel control samples. As expected, PepT1 was particularly abundant in intestinal cell lines and in the small bowel, whereas it was less abundant in the colon (Fig. 4B).

DISCUSSION

This study describes a morphological colonic adaptation in a homogenous series of patients with SBS type II (with a jejunocolonic anastomosis) compared with controls. The adaptive changes included a significant increase in crypt depth and number of cells per crypt. The remnant colon displayed an unchanged proliferative/apoptotic level and well-preserved absorptive NHE2, NHE3, and PepT1 transporter level. This is the first demonstration of controlled hyperplasia in human colonic epithelial cells of patients with SBS.

This series of patients with SBS appears to be representative of patients with SBS type II, with significant small bowel resection associated with severe malabsorption (31, 32). Our patients had good nutritional status with no inflammation, confirmed by normal blood CRP, and they displayed hyperphagia (2,600 kcal/day). Hyperphagia, (oral intake >1.5 times REE), has been demonstrated in human adults with postsurgical malabsorption (11, 12, 32). This oral behavior favors intestinal adaptation by inducing adaptive functional and morphological changes within the gut in animal models (5, 6, 8, 21, 29, 30). Patients also had a severe malabsorption with fewer malabsorbed carbohydrates ($88 \pm 11\%$) than proteins or lipids. On the basis of the hypothesis that adaptation occurs from 3 mo to 2 yr after bowel continuity was reestablished (35), we evaluated modifications in the remaining colon in hyperphagic patients whose surgery for anastomosis between the jejunum and the remaining colon had been performed at least 2 yr before.

In animal models, an increase in the mucosal mass of the colon has been extensively documented as an adaptive process (22, 34, 38, 53). The present study clearly shows adaptive morphological changes, with significantly deeper crypts

(+35%) and increased numbers of cells (+22%) per crypt, suggesting that the absorptive surface was increased in the colon after massive and distal small bowel resection. A previous study did not document such morphological changes in the colon of three patients with SBS with type II (jejunocolonic anastomosis), but the median time since constitution of these patients was less than 2 yr (50, 55). In animal models with small bowel resection, the adaptive response is temporally regulated that confirms the importance of studying time (15). Despite a deeper crypt depth, the ratio of proliferative/apoptotic cells to total cells was similar in patients and controls. On the basis of these data, we suggest that an adaptive hyperproliferative process leading to the formation of more cells in

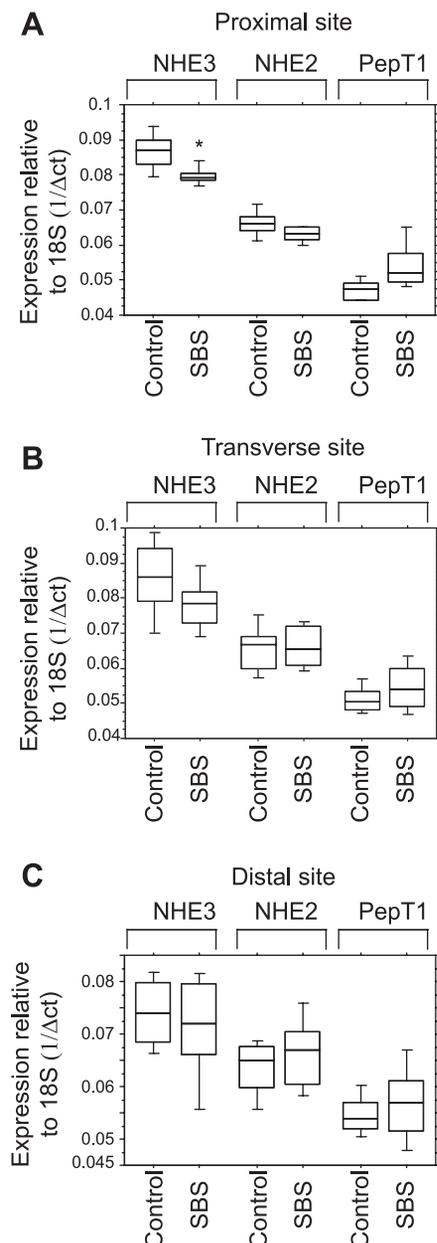


Fig. 3. Similar expression of NHE3, NHE2, and PepT1 mRNA in control group and patients with SBS. NHE3, NHE2, and PepT1 mRNA content was estimated by real-time RT-PCR and expressed as $1/\Delta\text{Ct}$ [$1/(\text{Ct target gene}-\text{Ct18s})$]. A: patients with SBS, $n = 6$ and controls, $n = 6$; * $P < 0.05$. B: SBS $n = 10$ and control group $n = 8$. C: SBS $n = 12$ and control group $n = 7$.

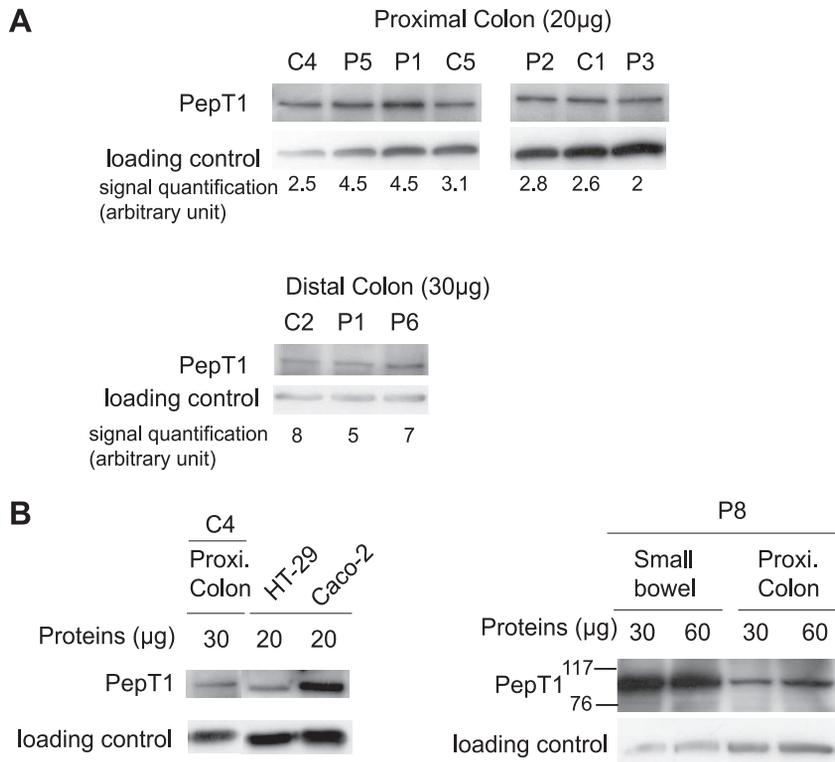


Fig. 4. Equivalent amount of PepT1 protein in patients and in controls. Gels for Western blot were loaded with 20, 30, or 60 μ g of total proteins as indicated. The membranes were successively hybridized with PepT1 and GAPDH (loading control) antibodies. *A*: PepT1 detection in biopsies from the proximal and distal sites of the colon from patients with SBS (P1, P2, P3, P5, and P6) and controls (C1, C2, C4, and C5). For each lane, the data represented the PepT1 signal divided by the signal of loading control and are expressed as arbitrary units. *B*: PepT1 detection in 2 cell lines (HT-29 and Caco-2) and in biopsies from small intestine (P8) and colon (P8 and C4). The weight markers indicated that the PepT1 band was detected between 76 and 117 kDa. Proxi, proximal.

deeper crypt occurred early after the resection and was then compensated with time. The maintenance of a controlled proliferating activity, which we observed at least 2 yr after the resection, is related to personal observation suggesting that patients with SBS have no predisposition to the development of colon cancer (prospective cohort of 22 patients with SBS, median age of 50.9 yr and a follow up to 29 yr).

During the past 10 yr, glucagon-like peptide (GLP)-2 has received attention as a therapeutic agent in the treatment of short bowel in humans. This hormone (secreted by the intestinal L cells in the ileum and the colon) displays pronounced intestinotrophic properties, by increasing mucosal growth, villus height, and water electrolyte absorption in animals (43). In humans with SBS, the administration of [glycine₂]-GLP-2 significantly increased in seven patients with SBS type I (end-jejunostomy) villus height, crypt depth, mitotic index, and intestinal function of the remnant small bowel (25). In a recent work, the confirmed enhancement of the small intestine and colon structure with a GLP-2 analog treatment was associated to an unchanged level of the protein content (49). After our demonstration of nonpharmacological colonic hyperplastic adaptation, a better understanding of its mechanisms deserves attention and may lead to the development of targeted treatments in patients with short bowel.

In rats, it has been suggested that colonic upregulation of NHE2 and NHE3 occurs as an adaptation process (34). However, our laboratory did not confirm this overexpression (28). Therefore, the few available data on NHE2 and NHE3 found in animal resection models remain conflicting. For the first time, the expression of NHE2 and NHE3 was studied in the colonic mucosa in humans. The NHE2 mRNA level remained unchanged in patients with type II compared with controls. The

NHE3 mRNA level was slightly decreased at the proximal site and equivalent to that of controls in the transverse and distal sites. This site-specific decrease close to the jejunocolonic anastomosis could be linked to local extracellular/intraluminal stimuli, such as modifications in osmolarity, luminal food-derived residues such as short chain fatty acids, and glucose or acidification (16, 42). Considering the fact that the decreased expression of NHE3 was weak and only observed near the anastomosis, we consider that NHE2 and NHE3 mRNAs remain stable in most of the remaining colon following major small bowel resection. Although the small intestine is the major site of PepT1 expression, it was also detected, to a lesser extent, in the colon (Figs. 2 and 4). Some authors have suggested that PepT1 enhanced expression in the colon is associated with the development of colon injury (40, 45, 46). Other authors have described PepT1 in the colon of unresected and resected rats (28) and resected and nonresected humans (17, 55). Our results provide additional support to the hypothesis that both mRNA but also protein PepT1 are present, though weakly, in the nondiseased human colon.

A previous study described an increase in PepT1 in patients with SBS (55), but in this work we did not observe such an upregulation of PepT1 at mRNA and protein levels. This apparent discrepancy could be linked to the diagnosis of pathology. In our cohort, we excluded patients with an inflammatory disease because PepT1 is involved in the intestinal inflammation activation pathway (9, 14, 52). None of five subjects (with a remaining colon) sampled by Ziegler et al. (55) had mesenteric ischemia. In addition, we considered only subjects at least 2 yr after resection, whereas Ziegler et al. (55) studied more acute response (several months after the last resection).

On the basis of the analysis of three transporters (the numbers of transporters was restrained because of a limited amount of biopsies samples), we suggest that the remnant colon keeps stable transport functions in the absence of small bowel. This is in accord with a recent study suggesting that ileal adaptation in a rat model occurs through cellular hyperplasia and not through cellular upregulation of transporters (24).

In conclusion, our study indicates that hyperphagic patients with SBS type II presented in the remaining colon a significant adaptive process characterized by an increased absorptive surface with an unchanged proliferative ratio and stable transporters amount. This is the first demonstration of controlled hyperplasia in human colonic epithelial cells of patients with SBS at least 2 yr after reestablishment of bowel continuity.

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