Bacterial DNA induces a proinflammatory immune response in patients with decompensated cirrhosis

We read with interest the study of Thalheimer et al (Gut 2005;54:556–63) in which they reviewed actual knowledge regarding the influence of infection on haemodynamics, variceal haemorrhage, hepatorenal effects, liver damage, and other effects.

We agree with these assumptions and would like to add information not quoted in the paper that may help explain some of the immune abnormalities usually found in patients with advanced decompensated cirrhosis. As the authors detailed in their paper, our group has reported on the detection of bacterial DNA in a significant proportion of patients with cirrhosis and culture negative non-neutrocytic ascites, and has also shown that these fragments may last in blood for variable periods of time.1 In our opinion, the presence of bacterial DNA is not only representative in itself of the presence of bacteria (either viable or non-viable) in our patients, but induces similar immunological changes as endotoxin or viable bacteria. The question of whether bacterial DNA also induces haemodynamic disturbances is currently under investigation.

Bacterial DNA contains a series of CpG motifs that join toll-like receptor 9 and activates a series of intracellular mechanisms leading to the synthesis of proinflammatory cytokines.2 We therefore observed that peritoneal white cells obtained from ascitic fluid in patients with the presence of bacterial DNA showed a marked activation pattern in the intracellular presence of cytokines involved in a type 1 immune response by means of flow cytometry was analysed,3 and also an increased ability to secrete these cytokines when cultured.4 Importantly, white cells in culture also displayed a significantly higher ability to secrete nitric oxide than cells obtained from patients without the presence of bacterial DNA, and nitric oxide levels showed a direct and significant relationship to the inducible form of nitric oxide synthase,5 suggesting that in this setting, as a consequence, nitric oxide synthesis is, at least in part, induced by this isoform.

Nitric oxide is a key agent in the pathogenesis of haemodynamic disturbances present in patients with advanced cirrhosis, and its levels are further increased in patients with hepaticorenal syndrome.6 Ascitic fluid nitric oxide levels are independently related to the development of renal impairment in patients with spontaneous bacterial peritonitis.7 Thus the relation between the presence of bacterial DNA in blood and the ability to secrete proinflammatory cytokines and nitric oxide by cells of the immune system in patients with decompensated cirrhosis suggest that endotoxin and viable bacteria should not only be taken into account in the design of new research protocols, but also bacterial DNA, or similar molecules, as demonstration of the presence of bacteria in patients with advanced cirrhosis.

Author's reply

We are grateful to Such et al for their comments on our review. As we had outlined, the influence of bacterial infection on the pathophysiology of cirrhosis is indeed an important one and Such et al have contributed significantly to this topic.8,9 We were aware of their data, but unfortunately some of it could not be retained in the final version of our paper due to editorial restrictions. Nevertheless, we agree that the presence of bacterial DNA, in the absence of viable bacteria or endotoxemia, might be an additional step in the sequence of events outlined in fig 2 of our review, even preliminary to endotoxemia.

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Conflict of interest: None declared.

References


Author's reply

We are grateful to Such et al for their comments on our review. As we had outlined, the influence of bacterial infection on the pathophysiology of cirrhosis is indeed an important one and Such et al have contributed significantly to this topic.8,9 We were aware of their data, but unfortunately some of it could not be retained in the final version of our paper due to editorial restrictions. Nevertheless, we agree that the presence of bacterial DNA, in the absence of viable bacteria or endotoxemia, might be an additional step in the sequence of events outlined in fig 2 of our review, even preliminary to endotoxemia.

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References


Perinatal passive smoke exposure may be more important than childhood exposure in the risk of developing childhood IBD

The large case control study of patients with inflammatory bowel disease (IBD) in the French paediatric population by Baron et al has clarified the role of well established genetic and environmental risk factors, as well as suggesting novel environmental risk factors (Gut 2005;54:337–41). However, we caution the authors on dismissal of the role of passive smoking in the smoking of IBD development in childhood. Our own data would suggest that analysing smoking data during pregnancy and at birth may be more important in the development of childhood IBD, rather than assessing smoking during childhood and at disease onset, as performed in this current study.

We have performed a case control study in South East Scotland of children with early onset IBD, matching cases of IBD diagnosed at less than 16 years of age with same sex and age (±1 year) controls attending the same general practice.1 In total, we matched 62 pairs of cases and controls, with a median age of disease onset in cases of 10.6 years. We demonstrated that parental smoking during pregnancy and around the time of birth was more common in parents of IBD cases, at 54% compared with control parents at 29% (p = 0.01; odds ratio (OR) 2.87 (95% confidence interval (CI) 1.05–16.97)). There was no significant effect seen when paternal smoking in pregnancy and at birth was analysed in IBD cases versus controls (p = 0.27). These

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data replicate the publication by Lashner and colleagues who studied 72 IBD cases and controls and found a similar relationship to smoking at birth—which was increased in children who later developed IBD in childhood (OR 3.02) and CD in childhood (OR 5.32). The authors of this study also demonstrated that maternal smoking at birth was important in the development of IBD and CD.

We agree with the findings of Baron et al that parental/pasive smoke exposure outside of the perinatal period, including at the time of IBD diagnosis, is not associated with the risk of developing IBD as a child.

The only factor which by smoke exposure during pregnancy and at birth leads to an increased risk of childhood IBD can only be a subject for speculation, but it is interesting to note a recent study has demonstrated chromosomal abnormalities in fetal epithelial cells in women who smoke during pregnancy.

In conclusion, our study agrees with previously published data to suggest a role between passive smoke exposure during pregnancy and at birth with the risk of childhood development of IBD. When assessing passive smoking in relation to childhood onset IBD, investigators should survey smoke exposure in the perinatal period and during childhood.

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Conflict of interest: None declared.

Reference

Author’s reply
We thank Russell et al for their interest in our study, concerning the link between passive smoking and the risk of IBD in children.

We agree that it is important to take into account the role of passive smoking not only during childhood and at disease onset but also during childhood. We also looked at this point in our study but came to different conclusions: 9.6% of mothers of IBD patients smoked during pregnancy versus 9.25% of control mothers (odds ratio (OR) 0.95 (95% confidence interval (CI) 0.53–1.72); p = 0.87). When considering only mothers of Crohn’s disease patients and control mothers, values were 9.9% and 9.5%, respectively (OR 0.95 (95% CI 0.50–1.81); p = 0.87). Moreover, concerning passive smoking and the risk of development of IBD, the findings were 14.2% and 12.8% for IBD patients and controls, respectively (OR 0.87 (95% CI 0.52–1.46); p = 0.60) and 15.3% for Crohn’s disease patients versus 14.4% for controls (OR 0.92 (95% CI 0.53–1.61); p = 0.77).

Due to the high number of questions and findings in our case control study, we only reported positive findings and what we considered as being the most important negative results. In conclusion, we confirm that in our study we could not link between IBD and passive smoking, including exposure during pregnancy and at birth.

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Conflict of interest: None declared.

An alternative to prophylactic colectomy for colon cancer prevention in HNPCC syndrome

The surgical option for treatment of a patient with screen detected colorectal cancer (CRC) from a family with hereditary non-polyposis colorectal cancer (HNPCC) is subtotal colectomy or segmental resection. Using decision analysis, we showed that subtotal colectomy performed at a young age leads to an increased life expectancy (LE) of 1–2.3 years. Based on these results and the high risk of developing a second CRC, we concluded that if CRC is detected in a young patient participating in a surveillance programme, colectomy with ileorectal anastomosis seems to be the treatment of choice.

A French Committee on HNPCC commented on our study. Firstly, they stated that using quality adjusted LE would be a more accurate approach for colorectal cancer patients. Secondly, the committee considered our five year survival rates optimistic. The five year survival rates for HNPCC patients with Dukes’ B cancer varied in the literature from 70% to 91% and those for patients with Dukes’ C from 19% to 70%.

These survival rates are lower than those used in our analysis. Thirdly, the committee mentioned that the overall five year survival of patients with CRC in HNPCC is approximately 55%. They stated that if the decision for an extended resection was made before the pathological staging of the tumour is known, 45% of patients will sustain a substantial decrease in QOL with no counterpart in quantity (that is, LE).

We agree that the patient’s choice is pivotal in decision making.

Due to the high number of questions and findings in our case control study, we only reported positive findings and what we considered as being the most important negative results. In conclusion, we confirm that in our study we could not link between IBD and passive smoking, including exposure during pregnancy and at birth.

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Defective denominators

I was interested in the paper by Langlands et al in which "prebiotic" carbohydrates altered the mucosal flora but apparently had no effect on cell proliferation (Gut 2004;53: 1610–16). The matter is of some importance as the products of in vivo fermentation (short chain fatty acids) may increase epithelial cell proliferation, leading to the possibility that such supplements could actually enhance the risk of colorectal cancer.1 2

The authors state that methodology (of gut microbiology) is always an important issue and I agree that this also applies to cell proliferation studies, as the results of the present work may be misleading on two counts. Firstly, I would never recommend the use of Ki67 as a marker of cell proliferation as: (1) the method is difficult to standardise; (2) the antigen has a long half life; and (3) anomalous expression has been demonstrated in non-cycling near tumours and after administration of growth factors. 3 For sections, Ki67 is far better however even using this antibody the results of the present study are unlikely to be conclusive as only 2–4 crypts could be scored. For most studies I would recommend scoring 30 hemi crypts.

The second point is that reliance on labelling indices can be misleading as lack of difference necessarily means no proliferative change as both sides of the ratio (labelled cells divided by number of cells) could have altered. This was demonstrated in our studies of epidermal growth factor in paracetamol fed rats where no differences in labelling index between orally fed and parenterally fed rats could be seen despite halving tissue weight and crypt cell production. When the data were re-expressed as labelling per crypt, the effects of treatment became apparent, a similar effect was seen in the stomach following misoprostol treatment.4

There is however a far easier and well validated method available for the study of human tissue. This is the so-called microdissection technique in which small pieces of stained material are teased apart and mitotic figures scored.5 This literally allows one to score over 100 crypts (if so wished) and as the results are expressed per crypt the effects of changes in denominator are automatically accounted for.

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Conflict of interest: None declared.

References


Author’s reply

We thank Goodlad for his interest in our article. In our study (Gut 2004;53:1610–16), we assessed expression of Ki67 in three markers most commonly used to indicate cell cycle entry in tissue sections. Importantly, there was no difference in the data obtained for all three. We agree that proliferating cell nuclear antigen is of limited value for the reasons mentioned by Goodlad and also the fact that the protein has a role in DNA repair, which reduces its specificity as a cell cycle marker. Similarly, Ki67 is not expressed by all cycling cells, may be downregulated by nutritional deprivation, and may also be involved in non-cell cycle related processes, such as ribosomal biosynthesis.2

We consider that the most useful markers of cycling cells are the minichromosome maintenance (MCM) proteins, which are abundant at all phases of the cell cycle and are downregulated following entry into quiescence, differentiation, or senescence.2 MCMs therefore provide a sensitive and specific indication of cell cycle entry. In our opinion these markers are preferable to counting mitotic figures, which is a subjective and error prone exercise that by definition provides a limited phase specific indication of cell cycle state in histological sections.

We agree that proliferation indices can be misleading and that when assessing large bowel crypts it is important to determine the number of labelled cells per crypt.3 We confirm that the mucosa in all subjects in our study was microscopically normal, as well as macroscopically normal, as stated. In particular, there was no difference in crypt length and number of cells per crypt between the study groups. The labelling indices determined were therefore valid indicators of cell cycle entry in the samples investigated.

Prebiotic carbohydrates, such as those used in our study, are completely fermented in the large bowel and none is excreted in faeces. The principal products of this fermentation are short chain fatty acids (SCFA). While SCFA have been associated with increased cell proliferation in some animal models, it is hard to believe that what are the major anions in the colon of all mammalian species should enhance the risk of cancer, particularly since one of these fatty acids, butyrate, is thought to be a differentiating agent. Fermented carbohydrates, such as dietary fibre, when measured properly in the diet, appear to protect against colorectal cancer in observational studies.4 The observed lack of effect of prebiotic carbohydrates on colonic crypt proliferation in our study suggests that a substantial increase in fermentable carbohydrate intake, as provided by these prebiotics, does not enhance proliferation, as shown in some animal models, and thus might be regarded as adding to the protective role of the fermentable non-starch polysaccharides (fibre).

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Conflict of interest: None declared.

References


Author’s reply

Further to Cummings and Coleman’s reply to my letter above, I would like to question the advocacy of minichromosome maintenance (MCM) proteins as proliferative markers, as the number of MCM positive cells can greatly exceed other labels and they are widely distributed on unreplicated chromatin. They would appear to be more of an indicator of replication potential and, as such, are likely to be useful markers of dysplasia.5 In addition, scoring immunohistochemical labelled cells is just as, if not more, "subjective and error prone" than scoring mitotic figures (which are far easier to score in "squash" preparations than in sections). My main
Interpreting observational and intervention studies of fibre has filled many journal pages in recent years. There are numerous problems which, in the context of the present discussion, relate primarily to people treating all sources of fibre as being equal, thinking that fibre supplements will have the same effect as fibre present in whole foods in the diet and the amounts of fibre considered to be protective. With regard to the study by Alberts and colleagues, the fibre was provided as a supplement and was only of wheat bran. As Goodlad and Alferes correctly note, the EPIC study showed a protective effect for fibre when intrinsically part of the diet, and from mixed sources. In other words, it is a high fibre diet that protects. The Bonithon-Kopp study used a fibre supplement, ispaghula, not found in most of the world, and at a very small dose of only about 3 g/day.

Author’s reply
We sought to identify cells at any point of the cell cycle, regardless of the rate of cycling or the duration of particular cell cycle phases. We therefore selected not to assess individual cell cycle phases in our samples, either by immunostaining or by counting mitotic figures. While additional roles for minichromosomes have been suggested, there is strong evidence that they function as essential replication factors. MCMs are displaced from chromatin following DNA replication, yet remain abundant in the nucleus throughout the cell cycle. Importantly, several groups have shown that MCMs are lost following cell cycle exit (into differentiation, quiescence, or senescence). MCMs are therefore useful immunohistochemical markers of cell cycle state. It is not surprising that MCMs are more abundant than Ki67 and proliferating cell nuclear antigen (PCNA), as the latter markers are not detectable in all cycling cells. Objectively, there is no difficulty in the interpretation of immunohistochemical staining when the marker used. Some markers, such as PCNA, produce substantial variation in staining intensity and cause difficulty in slide interpretation. However, our MCM antibodies have not provided us with such difficulties, resulting in low interobserver variation in numerous studies to date.

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References

Recurrence of exhausting hiccup in a patient treated with chemotherapy for metastatic colon cancer
A 61 year old man was surgically treated for a pT3 N1 G2 MO adenocarcinoma of the colon in February 2003. Immediately after surgery, an enteric fistula occurred that caused a delay in administration of adjuvant treatment. At the start of adjuvant chemotherapy (CT) in May 2003, CEA level was 18.2 ng/ml and a PET scan of the thorax and abdomen revealed the early appearance of two metastatic lesions in the liver. The patient underwent liver metastasectomy and in July 2003 was started on post-surgical systemic CT with the FOL-FIRI (leucovorin, 5-fluorouracil, oxaliplatin) regimen. After day 1 of CT, recurrence of an exhausting hiccup was observed that continued for nine days after therapy. No benefit from the re-use of chlorpromazine was obtained.

Notably, while undergoing the two CT regimens, the patient had received intravenous ondansetron (8 mg) plus intravenous dexamethasone (8 mg), which was used for prophylaxis of delayed emesis. In order to identify the causative drug of hiccup and taking into consideration previous reports indicating dexamethasone as a possible cause of hiccup, during the following cycles of CT this drug was omitted. This approach allowed the patient to continue CT without recurrence of hiccup.

This strong temporal relation between dexamethasone administration and occurrence of hiccup indicated that this drug was the cause of the patient’s hiccup. Moreover, discontinuing dexamethasone was sufficient to achieve disappearance of hiccup without any further pharmacological intervention.

The mechanism of corticosteroid induced hiccup is unknown, although some hypotheses have been proposed. For example, it has been suggested that there is a hiccup centre in the midbrain that receives input from the thoracic sympathetic nerves and the pharyngeal plexus. It has been proposed that stimulation of the midbrain or these various pathways may be responsible for production of hiccup. Moreover, animal studies suggested that corticosteroids may reduce the synaptic transmission threshold in the midbrain and affect the metabolism of brain neurotransmitters.

We reported our case to make oncologists aware that a symptom appearing during CT treatment (hiccup in our case) should not always be ascribed to the use of antineoplastic drugs. It is also true that some cytotoxic drugs, such as irinotecan and oxaliplatin, have been implicated as a cause of hiccup. In particular, the incidence of hiccup after treatment with irinotecan was reported in 49/16518 patients and, for other cytotoxic drugs, almost exclusively in men (49/9313).

Conflict of interest: None declared.
Laterally spreading tumour in which interstitial deletion of β-catenin exon 3 was detected

Laterally spreading tumours (LSTs) of the colon and rectum are defined as lesions greater than 10 mm in diameter with a low vertical axis that extend laterally along the luminal wall. As most LSTs remain as adenomas or early invasive cancers, LSTs have been thought to have relatively little malignant potential. LSTs are divided into two macroscopic subtypes: flat (F)-type, which is composed of superficially spreading lesions with flat and smooth surfaces, and granular (G)-type, which is composed of superficially spreading aggregates of nodules. Despite distinctive biological behaviours of LSTs, only a few genetic alterations have been reported, such as K-ras and p53 mutations and cyclooxygenase 2 overexpression. A 62 year old Japanese woman was referred to our hospital for treatment of a colonic tumour. Colonoscopy in our hospital showed an F-type LST with a central depression surrounded by a flat elevated area in the caecum (fig 1A). Microscopically, the tumour consisted of a well differentiated adenocarcinoma with a tubular adenoma and had invaded the submucosal layer.

After obtaining informed consent from the patient, genetic analysis was carried out. No genetic alterations were found in APC, K-ras, or p53 genes. To clarify relevant alterations of gene expression, we analysed the gene expression profiles by a cDNA array. Among 350 cancer related genes, bone morphogenic protein 4 (BMP4) was one of the most differentially expressed genes in tumour tissues and matched normal tissues (fig 1B). BMP4 is a member of the transforming growth factor superfamily of growth factors. As BMP4 expression is reportedly correlated with onco- genic β-catenin in human colon cancer cells, we analysed alterations in β-catenin in tumour tissues. Intense nuclear expression of β-catenin was immunohistochemically seen within the nuclei of tumour cells. (D) Intestinal deletion examined by polymerase chain reaction spanning the genomic region flanking exon 3 and the surrounding introns. A shorter band was detected in both carcinoma and adenoma tissues compared with the normal size of 931 bp. CA, carcinoma tissue; TA, tubular adenoma tissue; N, normal tissue.

Figure 1 (A) Endoscopic picture with indigocarmine dye spraying showing an F-type laterally spreading tumour with a central depression surrounded by a flat elevated area in the caecum. (B) cDNA array hybridisation image of the tumour and non-tumour tissues. bone morphogenic protein 4 (BMP4) was one of the most differentially expressed genes in the tumour tissues and matched normal tissues. (C) Intense nuclear expression of β-catenin immunohistochemically seen within the nuclei of tumour cells. (D) Intestinal deletion examined by polymerase chain reaction spanning the genomic region flanking exon 3 and the surrounding introns. A shorter band was detected in both carcinoma and adenoma tissues compared with the normal size of 931 bp. CA, carcinoma tissue; TA, tubular adenoma tissue; N, normal tissue.

![Image](https://www.gutjnl.com/)

Figure 2 DNA sequencing showing interstitial deletion of the 394 bp region in tumor tissue. Three base inverted repeats, AGC and GCT, were found in sequences flanking the interstitial deletion. The deletion included the part of exon 3 containing critical serine and threonine codons for GSK-3β phosphorylation.

![Image](https://www.gutjnl.com/)
present patient had no past history or family history of cancer. It would be interesting to investigate whether β-catenin mutation positive HNPCC cancers have any specific morphological features.

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References

Functional role of the 503F variant of the organic cation transporter OCTN1 in Crohn’s disease

Several susceptible gene loci were identified as being involved in the etiology of Crohn’s disease (CD).1 Recently, a non-synonymous single nucleotide polymorphism in the SLC22A4 gene encoding the organic cation transporter OCTN1 has been linked with CD in Caucasian populations (a 1672CT transversion, resulting in the amino acid substitution L503F).2,3 However, the functional consequences of this alteration are unclear as yet.

We have now discovered that L-ergothioneine (ET, 2-mercaptophistidine trimethylbainaine), a naturally occurring water soluble thiol compound of dietary origin, is a physiological substrate of OCTN1.4 Analysis of the concentration dependence of ET transport in OCTN1 transfected HEK293 fibroblasts by liquid chromatography tandem mass spectrometry revealed that the 503F variant was associated with a threefold higher substrate affinity (1/Km) and a twofold lower maximal transport velocity (Vmax), which resulted in a 50% higher initial transport capacity (Vmax/Km (503F) = 1.5 × Vmax/Km (503L)) at low ET levels (<10 μmol/l) (fig 1A). Analysis of the time course of ET transport showed a higher clearance for the 503F variant (CL (503F) = 1.65 × CL (503L) at an ET concentration of 10 μmol/l) (fig 1B). ET transport by 503L and 503F was sodium

Figure 1 Ergothioneine and OCTN1. Concentration dependence, Km, and Vmax of specific ergothioneine (ET) uptake in HEK293 cells constitutively expressing the 503L variant or the 503F variant after one minute of loading (A); specific uptake and clearance (CL) over a time course after incubation with 10 μmol/l ET (B); effects of sodium (C) and pH (D) on specific uptake after one minute of loading with 10 μmol/l ET. In sodium reduced transport buffer, NaCl was isotonically replaced with choline chloride (which did not interfere with ET transport). An equal expression level of both OCTN1 mRNAs was controlled by quantitative real time polymerase chain reaction (TaqMan assay). Linear correlation of ET concentrations in CD14+ monocytes (fractionated from peripheral blood mononuclear cells) with OCTN1 mRNA expression (relative to the housekeeping gene GAPDH, lowest expression was set to 1) in eight healthy volunteers that were homozygous carriers of the 503L variant (E). MTT assay7 of the proliferation of Caco-2 colon tumour cells with and without OCTN1 mRNA expression after 24 hours of incubation with ET or glutathione. Resulting formazan formation was determined by absorbance at 568 nm (F). Data are means (SEM) of three (A–D) or 8–16 (F) independent experiments. *p<0.05, **p<0.01, ***p<0.001: significant differences between OCTN1 variants (A–D); significant differences compared with buffer controls (F), as determined by one way ANOVA with Holm-Sidak correction (α=0.05).

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and pH dependent; only at unphysiologically low Na⁺ and pH values were the differences in transport activity between both variants lost (fig 1C). Considering that maximal levels of ET found in tissues and in common foods are in the nanomolar to low micromolar range, our data suggest that carriers of the 503F allele accumulate higher ET concentrations in OCTN1 expressing cells compared with carriers of the wild-type 503L allele. Therefore, high tissue levels of ET may constitute a possible risk factor for CD.

The involvement of OCTN1 in the inflammatory process is further supported by observations that OCTN1 is strongly expressed in intestinal epithelial and immunological cells, particularly in CD4⁺ monocytes/macrophages playing a key role in the immunopathogenesis of CD, as well as by the finding that levels of SLC22A4 mRNA were upregulated by proinflammatory cytokines such as tumour necrosis factor α. Moreover, we found transcriptional regulation of SLC22A4 to determine essentially ET uptake: in CD4⁺ monocytes lacking the 503L variant OCTN1 expression, expression levels of SLC22A4 mRNA showed high interindividual heterogeneity and were directly proportional to cellular ET content (fig 1H). Accordingly, in CD4⁺ and CD8⁺ lymphocytes lacking OCTN1 expression, we detected no ET (data not shown).

The physiological or pathophysiological functions of ET are as yet unknown. We tested the effect of ET on proliferation of the colon cancer epithelial cell line Caco-2 that was shown to be homozygous for the susceptible 503F allele and to express high levels of OCTN1 mRNA. Cell proliferation was enhanced in a dose dependent manner after exposure to ET concentrations above 20 μmol/l for 24 hours: at 200 μmol/l, proliferation was increased to 120 (3)% of the control. In contrast, stimulation of proliferation was seen when a Caco-2 variant without OCTN1 expression was employed; consequently, after treatment with 200 μmol/l ET, only diffusion correct OCTN1 uptake to 0.67 (0.03) pmol/mg protein occurred. When incubated with gluthione, both Caco-2 cell lines exhibited an antioxidant typical inhibition of proliferation that was independent of OCTN1 expression (fig 1I). ET, therefore, rather than antioxidant activities, stimulatory effects on cell proliferation appear to constitute the functional role of ET. ET may accelerate the inflammatory process by transcriptional activation of fibroblast repair proliferation, thereby also conferring susceptibility of CD patients to develop colorectal cancer.

Collectively, our data suggest that the OCTN1 substrate ET is a proliferative factor in inflammatory diseases such as CD, and subjects carrying the 503F allele are at an increased risk due to a higher intracellular accumulation of ET.

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Diarrhoea as a presentation of bird flu infection: a summary on its correlation to outcome in Thai cases

Bird flu or avian flu, caused by H5N1 virus, is a new emerging infectious disease. There has been worldwide avian influenza infections in poultry since 1997. Recently, H5N1 caused severe disease with high mortality in humans in Vietnam and Thailand. Most infected cases usually developed progressive pneumonia with acute respiratory distress syndrome and consequently died. Antipodal presentations of patients with bird flu were also noted. Leong et al recently reported a fatal bird flu infected case in Vietnam with a presentation of diarrhoea, without respiratory symptoms. I performed a mini-study in order to document the magnitude of diarrhoeal presentation among reported Thai patients and the correlation with outcome. A literature review on papers concerning human bird flu in Thailand was performed using databases of published works cited in Index Medicus and the Science Citation Index. I also reviewed published local and 256 local Thai journals, which are not included in the international citation index, for reports of human bird flu infection in Thailand. Studies that contained incomplete data were excluded from further analysis.

Six reports1-5 of 12 Thai patients with a confirmed diagnosis of bird flu were found. Of 12 infected cases, respiratory symptoms were seen in all cases and diarrhoea was detected at presentation in five cases (41.7%). Considering the five diarrhoeal cases, acute disease and development of ARDS was detected in four cases and there were three deaths. Concerning the seven non-diarrhoeal cases, ARDS was detected in five cases and there were five fatalities. There was no significant correlation between presentation of diarrhoea and development of ARDS (p = 0.05) or mortality (p = 0.05) but there was a significant correlation between the development of ARDS and mortality (p = 0.001). There are some reports of diarrhoea in severe bird flu infection. Poovorawan recently proposed that diarrhoea was an important presentation of bird flu and could imply a poor prognosis. Here, I attempted to assess the magnitude of diarrhoea among Thai infected cases, and its correlation with infection outcome. According to this study, the prevalence of diarrhoeal presentation was high, similar to a recent study in Vietnam (approximately 70%). Therefore I conclude that diarrhoeal presentation had a poor correlation with outcome among our subjects.

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References


High levels of disease related prion protein in the ileum in variant Creutzfeldt-Jakob disease

Disease related prion protein (PrPSc) is readily detectable in lymphoreticular tissues in variant Creutzfeldt-Jakob disease (vCJD) but not in other forms of human prion disease. This distinctive pathogenesis together with the unknown population prevalence of asymptomatic vCJD infection1-4 has led to significant
concerns that secondary transmission of vCJD prions will occur through a wide range of surgical procedures. Risk assessment for intestinal endoscopy, biopsy, and surgery is currently limited by a lack of knowledge about relative PrPSc levels and prion titres within intestinal tissues in vCJD patients. Because of its high content of lymphoid follicles, terminal ileum is regarded as the intestinal tissue having the highest potential for iatrogenic transmission of vCJD prions. Here we provide the first report of relative PrPSc concentrations in vCJD terminal ileum.

Tissues were obtained at autopsy with consent from relatives from four patients with neuropathologically confirmed vCJD and two patients with neuropathologically confirmed sporadic CJD (both PRNP codon 129MM with type 2 PrPSc in brain). Terminal ileum was analysed for PrPSc by high sensitivity immunoblotting and for abnormal PrP immunoreactivity by immunohistochemistry. Using these methods, terminal ileum from all four vCJD cases showed high levels of detectable PrPSc (fig 1A). In three vCJD cases, 2/2 homogenates prepared from each ileum specimen were positive for PrPSc whereas 2/4 ileum homogenates were positive in the other vCJD case. The glycoform ratio of protease resistant fragments of di- mono-, and non-glycosylated PrP in terminal ileum appeared to be closely similar to the type 4 PrPSc pattern seen in vCJD tonsil.

Although there was variation in PrPSc concentration between different homogenates of vCJD terminal ileum, PrPSc levels in positive samples were typically in the range 0.1–1% of that present in vCJD brain (fig 1B). With respect to both sampling variation and PrPSc concentration, terminal ileum appears to be closely similar to lymph nodes in vCJD. These findings, together with our previous studies, show that PrPSc deposition within the intestine is not uniform in vCJD. From the four cases of vCJD with PrPSc positive terminal ileum studied here, 0/2 cases with available tissue had detectable PrPSc in the appendix and only 1/3 cases had detectable PrPSc in the rectum. In contrast with findings in vCJD terminal ileum, no detectable PrPSc was found in homogenates of terminal ileum prepared from sporadic CJD patients (fig 1A). The lack of detection of PrPSc in sporadic CJD terminal ileum extends our previous findings for one of these cases in which we have previously reported a lack of detectable PrPSc in tonsil, rectum, and appendix.

In agreement with findings from immunoblotting, immunohistochemistry showed abnormal PrP deposition in the terminal ileum in vCJD (fig 1C) but not in sporadic CJD (data not shown). The irregular distribution of abnormal PrP positive lymphoid follicles seen in vCJD terminal ileum is consistent with variation in PrPSc concentration detected in different terminal ileum samples by immunoblotting.

Albeit from necessarily limited numbers investigated, the uniform presence of PrPSc in vCJD terminal ileum, at concentrations of up to 1% of those found in vCJD brain, reinforces concerns that iatrogenic transmission of vCJD prions might occur through contaminated intestinal endoscopes, biopsy forceps, or surgical instruments. These findings should assist policy makers in the UK and elsewhere in risk assessments about the use of disposable forceps for intestinal biopsy. Alternative approaches to risk reduction may now be possible as practical means of prion decontamination for endoscopes and surgical instruments are now feasible using enzymatic methods.

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Conflict of interest: declared (the declaration can be viewed on the Gut website at http://www.gutjnl.com/supplemental)

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3 Wadsworth JDF, Joiner S, Hill AF, et al. Tissue distribution of protease resistant prion protein in variant CJD using a highly sensitive


Chronic inflammatory intestinal diseases and bone loss

We were very interested in the recent article by Moschen et al. on activation of the RANKL/OPG system in inflammatory bowel disease (IBD) (Gut 2005; 54:479–87). Until recently, osteoporosis secondary to gastrointestinal diseases was mainly considered a direct consequence of malabsorption. The article of Moschen et al. and a previous one of our group on bone loss in coeliac disease, a disorder similarly characterised by intestinal inflammation, offer a new perspective on the pathogenesis of bone loss and reveal a more complex picture. Moschen et al. demonstrated overproduction of OPG in the cells of colonic mucosa in IBD whereas Taranta and colleagues showed the direct role of the soluble cytokines in the serum of coeliac patients on bone cells. In fact, they found an increased RANKL/OPG ratio in untreated coeliac patients and different effects of the sera of untreated coeliac patients with respect to those on a gluten free diet, on cultured bone cells. These effects included increased in vitro osteoclastogenesis, and lower interleukin 18 and OPG expression in osteoblasts. In both studies, these biochemical observations were translated in a reduction of bone mass. Moschen et al. found a negative correlation between OPG plasma levels and spine and femoral z scores and bone mineral density (BM1). Taranta and colleagues observed a significant negative correlation between BM1 z score and interleukin 6 levels and RANKL/OPG ratio. In the discussion, Moschen et al. observed that “studies of OPG/RANKL and BM1 are required to validate” their model.

We believe that our study may be a first step towards understand, at least in part, the relationship of inflammation to bone loss in intestinal diseases. These results are also in accordance with recent studies on primary osteoporosis, which are beginning to show a relevant role of local and systemic factors on cell activity. Finally, these studies may also open the way to different therapeutic approaches—namely, drugs specifically acting on cytokines release and/or activity—for bone loss secondary to “inflammatory intestinal diseases”.

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References


Acid Related Diseases: Biology and Treatment


This textbook by Irvin Modlin and George Sachs is a welcome addition to the increasingly important and dynamic field of gastroenterology. The authors do provide a list of suggested reading for further introduction to the scientific literature. The information is generally presented in a refreshing and amicable style. I think the book is friendly enough to be of benefit to an average student, but at the same time it caters adequately for the more seasoned learner too. It features some beautiful pictures and drawings depicting many individuals who have contributed to this field over the last hundred or so years. I thought the cartoons in the chapter on Helicobacter pylori were particularly pleasing and informative.

I particularly liked the background given the development of the first proton pump inhibitor (PPI). I thought this was thoughtfully stimulating and will no doubt enable me to create a greater impression in front of the next PPI rep that I meet. The chapter on peptic ulcer disease is by and large par for the course, but the section on Barrett’s oesophagus presents a very logical and sensible approach towards tackling an area which remains controversial.

As a matter of personal taste, I would like to see a few key messages or take home points at the end of each chapter. These can also act as a quick source of reference for those who find that spare time is generally an elusive commodity, which, I suspect, is nearly all of us.

All in all, it is a timely and a creditable addition covering a very important and rapidly evolving field of gastroenterology, and the authors ought to be congratulated for their efforts. Would I buy it? Probably yes, but only if I did not have a copy of the first edition. I would certainly recommend it as a departmental book as, among its many virtues, it provides useful tibits to amuse the audience during presentations.

A Mahmood

BOOK REVIEW

Acid Related Diseases: Biology and Treatment


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CORRECTIONS

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In the August issue of Gut one of the authors was omitted from the paper by Goulding et al (G Goulding, A Murphy, G MacDonald, S Barrett, J Crowe, J Hegarty, S McKierman, and D Kelleher. The CCR5-A32 mutation: impact on disease outcome in individuals with hepatitis C infection from a single source. Gut 2005; 54:1157–1161). R McManus (Department of Clinical Medicine and the Dublin Molecular Medicine Centre, Trinity Centre for Health Sciences, St James Hospital, Dublin, Ireland) should have been listed as the second author on the paper.

In the August issue of Gut the following paper, Randomised controlled trial comparing percutaneous radiofrequency thermal ablation, percutaneous ethanol injection, and percutaneous acetic acid injection to treat hepatocellular carcinoma of 3 cm or less (S-M Lin, C-J Lin, C-C Lin, C-W Hsu, and Y-C Chen. Gut 2005; 54:1151–1156), was published without one of the author corrections being made. On page 1154 under the heading “Local and novel HCC recurrence”, the first line reads “…a median of 35 months” and should have been revised to “…a median of 24.3 months”.

Michael战备侦察兵Yong

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