Oesophageal cancer and gastro-oesophageal reflux: what is the relationship?

J Lagergren

There seems to be a causal link between reflux and oesophageal adenocarcinoma, and Barrett’s oesophagus is very likely an important, and possibly necessary, link in the causal pathway from reflux to oesophageal adenocarcinoma

The direct relation between reflux and adenocarcinoma of the oesophagus has recently been addressed in several well designed population based epidemiological studies, using different study designs. The studies are all in agreement that reflux is associated with an increased risk of oesophageal adenocarcinoma in a seemingly dose dependent manner.

The specific role of Barrett’s oesophagus for this association was not addressed in these studies but the presence of Barrett’s oesophagus has been regarded as a potential link in the causal pathway. More speculatively, based on an interpretation of the results from a large population based study in Sweden, we hypothesised that the relation between reflux and oesophageal adenocarcinoma might be independent of Barrett’s oesophagus. This hypothesis was based on two findings of our study. Firstly, among patients with a history of severe and longstanding reflux symptoms, the association with adenocarcinoma was of similar strength as that of the association previously reported between the occurrence of Barrett’s oesophagus and the risk of oesophageal adenocarcinoma in large scale studies. Secondly, when we compared our case patients with oesophageal adenocarcinoma who had a verified Barrett’s oesophagus with those who had no such mucosa detected during endoscopy or surgery, we found that the association between reflux and oesophageal adenocarcinoma was equally strong. The possibility of tumour overgrowth covering the Barrett might explain this finding. However, nevertheless, the hypothesis deserves further attention.

In this issue of Gut, Solaymani-Dodaran and colleagues address the role of Barrett’s oesophagus compared with the role of reflux per se in the aetiology of oesophageal adenocarcinoma in an interesting investigation [see page 1070]. They analysed the strength of the association with adenocarcinoma in three cohorts of patients, including a cohort of patients with Barrett’s oesophagus, a cohort of patients with oesophagitis, and third cohort of patients representing simple reflux. As a reference, a comparison cohort of patients without any of these conditions was selected. The point estimate of the relative risk of oesophageal cancer was 10 in the Barrett’s cohort while the corresponding risk estimates were approximately 2 in the two cohorts representing reflux without Barrett’s metaplasia. The authors conclude that their findings do not support the fact that gastro-oesophageal reflux in itself predisposes to oesophageal cancer. This interpretation is not necessarily true, however. There are some problems that deserve consideration in their study.

Firstly, the study included oesophageal cancers of any histological type in the analysis (that is, it did not specifically address the risk of adenocarcinoma of the oesophagus, the only type of oesophageal cancer that has been linked with reflux). Secondly, the risk of chance errors was high as the study, like most other prospective studies of this relatively rare disease, was underpowered. The limited number of 43 cases of oesophageal cancer were evaluated, and among these cancers the oesophageal squamous cell carcinomas and any other types were included. Thirdly, selection bias in the comparisons between the cohorts might explain the differences between the three cohorts reported in the study. The cohort representing Barrett’s oesophagus is likely to consist of patients with the most severe and longstanding reflux disease compared with the two comparison cohorts representing oesophagitis and simple reflux. From previous studies, we know that the relationship between reflux and oesophageal adenocarcinoma is strongly dose dependent both with regard to severity and duration of reflux (the relative risk of this cancer was 43.5 among patients with long term and severe symptoms in our Swedish investigation). We also know that patients who develop Barrett’s oesophagus usually have had reflux symptoms for long periods of time, typically for decades. The other two cohorts in the study by Solaymani-Dodaran and colleagues are likely to encompass those with, on average, less severe and less longstanding reflux. Any differences between reflux cohorts in the study are likely to be strongly affected by such selection bias. Hence the study cannot reliably verify whether or not Barrett’s oesophagus is a mandatory step in the development of adenocarcinoma of the oesophagus.

In summary, based on the current knowledge, there seems to be a causal link between reflux and oesophageal adenocarcinoma. Barrett’s oesophagus is very likely an important, and possibly necessary, link in the causal pathway from reflux to oesophageal adenocarcinoma but more studies are needed before the true answer to this important and interesting question can be established.


Correspondence to: Professor J Lagergren, Department of Surgery, Karolinska University Hospital, Stockholm SE-171 76, Sweden; jesper.lagergren@kus.se

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Coeliac disease

T helper cell polarisation in coeliac disease: any (T-)bet?

M H Holtmann, M F Neurath

Recent data strongly support the view that coeliac disease is a Th1 mediated inflammatory disease as both interferon \( \gamma \) production and T-bet levels in gut infiltrating cells are upregulated

The puzzling observation on high interferon \( \gamma \) (IFN-\( \gamma \)) but low interleukin (IL)-12 levels in coeliac disease (CD) has resulted in questions about the underlying principles of T helper cell polarisation. In this issue of Gut, the molecular basis of T helper cell polarisation in CD has been illuminated by the finding that T-bet, the master transcription factor of T helper cell type 1 (Th1) cells, is upregulated in this disease [see page 1090].

The past decade has witnessed a dramatic improvement in our pathophysiological understanding of inflammatory disorders of the intestine due to extensive research efforts focusing on regulatory mechanisms of the immune system. This has been greatly facilitated by the advent of molecular biological techniques which have allowed for the identification and characterisation of distinct factors in the complex regulatory network of the immune system. General themes have thus been recognised and an attempt is being made to classify diseases according to common immunological features.

One crucial emerging theme is the Th1/Th2 polarisation of T helper cells. On activation with distinct immunological stimuli, naïve CD4+ precursor T helper cells differentiate into mature T cells which either produce a so-called Th1 cytokine profile characterised by tumour necrosis factor \( \alpha \) (TNF-\( \alpha \)) and IFN-\( \gamma \), or a T helper cell type 2 (Th2) cytokine profile including IL-4, IL-5, IL-9, and IL-13. While these T cell subsets produce proinflammatory cytokines that can cause substantial tissue injury in vivo, more recent data suggest that additional subsets of CD4+ T helper cells exist with anti-inflammatory functions. These subsets include T helper 3 (Th3) cells producing transforming growth factor \( \beta \) (TGF-\( \beta \)), Tr1 cells producing IL-10, and CD25+CD4+ regulatory T cells (Treg) expressing the master transcription factor Foxp3.

Like all mucosal surfaces, the intestinal mucosa possesses a special immune system (gut associated lymphoid tissue (GALT)) that can be considered the biggest immune system of the body due to its large surface of several hundred square metres of human gut. The immune system of the upper gastrointestinal tract is physiologically exposed to a large amount of antigens from food, such as gluten, which is the protein fraction of wheat (here called gliadin), rye, and barley, and causes the stickiness of these cereals thus allowing the baking of bread.

Normally, ingestion of food antigens such as gliadin does not elicit an immune response. Otherwise, this would be detrimental because it would lead to a permanent state of inflammation of the bowel considering the large amount of food antigens. The intestinal immune system in general is characterised by a hyporesponsive state, which should not be mistaken for passive toleration, but should rather be considered as active inhibition of intestinal immune responses. Physiologically, an anti-inflammatory cytokine milieu prevails in the intestinal immune system with a predominance of Th3 and Treg T cells that contribute, together with other factors, to this hyporesponsive state. In CD, this hyporesponsiveness to gliadin is abrogated. While the initial pathogenetic steps have been fairly well characterised, details of the downstream immune response involving specific B and T cell responses are still a matter of debate.

The first report on a malabsorption syndrome on ingestion of wheat that resembles CD dates back to Aretaeus from Cappadocchia, 2nd century AD. In 1888, Gee gave the first detailed clinical description of the “coeliac affection”.

Today, CD can be considered as a model disease with autoimmune characteristics for several reasons. In contrast with most other autoimmune disorders, in CD an unequivocal trigger (the cereal protein gliadin) has been identified. The antigenicity can be related to a 33mer epitope that is generated by digestion with intestinal enzymes. In addition, there is an established close genetic association with the HLA haplotypes DQ2, found in 95% of patients, or DQ8 found in most of the remaining patients. A highly specific humoral autoimmune response with autoantibodies (IgA anti-endomysial antibodies), with almost 100% sensitivity and specificity to endogenous tissue transglutaminase (tTG), is elicited.

tTG is a calcium dependent ubiquitous enzyme that cross links matrix proteins and thus stabilises connective tissue. Gliadins, which are glutenin rich (>30%), are preferred substrates for tTG as donor substrates and substrate specificity is particularly high for the recently identified 33mer key epitope. Deamidation leads to gliadin-gliadin cross links and glutenase transfer to proteins in the extracellular matrix which probably generates additional antigenic epitopes. In addition, in the absence of glutamyl acceptor proteins, tTG can deamidate glutenase residues thus converting them into negatively charged glutamic acid. These modified gliadin peptides have a higher affinity to the coeliac specific HLA-DQ2 molecules than their non-deamidated precursors and can thus elicit a much stronger specific immune response.

A recent model proposes that autoreactive B cell clones specific for tTG or

References


Biopsies were taken from the distal intestine of untreated patients with CD. Production of IFN-γ was more than 1000-fold increased, consistent with a Th1 cytokine response, while IL-4 levels were below the level of detection. In patients treated with a gluten free diet, IFN-γ production was as low as in healthy controls but could be stimulated in vitro by gluten up to levels of untreated patients. Release of IFN-γ is most likely pathogenetically relevant as neutralisation of IFN-γ prevented gluten-mediated morphological changes in an ex vivo organ culture of treated CD epithelial cells and for development of another CD as a Th1-mediated disease because IL-12 cast doubt on the classification of CD as a Th1-mediated inflammatory disease as both T-box transcription factors GATA-3 and STAT-6 were upregulated in the mucosa of patients with untreated CD, but not from healthy controls, led to upregulation of T-bet. This upregulation of T-bet was mediated by STAT-1 and could be blocked by inhibition of STAT-1. T-bet is a strong regulator of IFN-γ production. Of note, it seems that conversely, expression of T-bet can also be regulated by IFN-γ mediated via STAT-1. This is interesting as T-bet acts at a very early stage in the regulatory pathway of Th1 commitment of T cells while IFN-γ production represents the functional phenotype of these cells.

The authors point to the limitation of their work which is that analysis of transcription factors was performed in whole biopsies and not in purified cell types. This might raise the concern that upregulation of T-bet could reflect increased infiltration of the mucosa by mononuclear cells rather than a true upregulation of T-bet in these cells. However, the fact that STAT-4 is not upregulated speaks against this explanation. Nevertheless, these findings are interesting. In fact, these data strongly support the view that CD is a Th1 mediated inflammatory disease as both IFN-γ production and T-bet levels in gut infiltrating cells are upregulated. The lack of STAT-4 upregulation fits to the reported absence of IL-12 in CD (see above). It thus seems that there is an alternative regulatory pathway for Th1 differentiation that starts from activation of T-bet but then bypasses STAT-4 and IL-12. It is thus tempting to speculate that T-bet activation in CD is induced or maintained by other cytokines beyond IL-12, such as IL-23 or...
IL-27. Further studies are necessary to address this point.

What is the relevance of this work? Analysis of pathomechanisms at the molecular level provides the potential for novel effective treatment strategies. The clear cut classification of Crohn’s disease as a Th1 mediated disease with the elucidation of the pivotal pathogenetic role of TNF-α has led to the development of innovative anti-TNF strategies by recombinant monoclonal antibodies. In CD, there is little need for effective anti-inflammatory treatment, however, as elimination of the exogenous trigger by a gluten free diet may lead to clinical, histomorphological, and immunological recovery.

Other questions are clinically more relevant. There must be variable environmental factors to explain why clinical manifestations can range from asymptomatic to severe malabsorption. CD is associated with a high prevalence of various other autoimmune disorders such as type 1 diabetes, dermatitis herpetiformis, autoimmune thyroiditis, collagen diseases, autoimmune alopeica, and autoimmune hepatitis. In 35% of patients in whom CD is diagnosed at age 20 years or older, including subclinical forms, these associated autoimmune disorders can be found. In patients who have been treated properly with a gluten free diet from infancy, the prevalence is only 5%. Clinically, there may be a need for effective screening methods to detect patients with subclinical CD.

Assignment of CD to the Th1 mediated diseases is of great scientific interest. The unique features of CD as a model autoimmune disease will facilitate clinical investigation of more general mechanisms of Th1 mediated diseases. The Th1/Th2 paradigm may, like all paradigms in science, be only temporary, and will probably give way to novel paradigms as research and science progress in the future. For time being, however, the Th1/Th2 paradigm is helpful in defining immunological disorders with the prospect of developing effective novel treatment approaches.


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Irritable bowel syndrome

East meets West: infection, nerves, and mast cells in the irritable bowel syndrome

S M Collins, G Barbara

The incidence of irritable bowel syndrome following acute gastroenteritis in China is similar to that reported in the UK and North America

The irritable bowel syndrome (IBS) is the most common gastroenterological disorder seen in Western societies where it imposes a major socioeconomic burden. The economic impact of IBS arises, at least in part, from difficulties in diagnosis and treatment, which in turn reflect our limited conceptualisation of this disorder. Recent studies have identified acute bacterial gastroenteritis as a strong risk factor for the development of IBS but evidence supporting this relationship is limited to studies performed in Western countries.

Although IBS has received less attention in non-Western countries, available data suggest that it is common in the African continent, Malaysia, and China where the condition is significantly underdiagnosed by Western practitioners. This hesitation in diagnosing IBS in Eastern countries likely reflects uncertainty regarding the influence of culture and endemic infection on the clinical expression of IBS in these countries. Indeed, in countries where enteric parasitic infestation is common, there is controversy regarding the role of infection in the development of IBS, as eradication of the parasite may not improve IBS symptoms. However, studies in animal models suggest that the host response to the initial infection is a critical determinant of persistent gut dysfunction, and the continued presence of the parasite is not necessary for induction of a chronic dysfunctional state. Thus the role of infection in the pathogenesis of IBS in these countries may be underestimated.

It is on this background of uncertainty that the paper by Wang and colleagues, published in this issue of Gut, is received with considerable interest [see page 1096]. This cohort study evaluated gastrointestinal symptoms in 295 patients recovering from acute gastroenteritis in Beijing, China. During a 1–2 year follow up period, the authors found an 8.1% incidence of IBS among all patients compared with 0.8% of controls, and an incidence of 10.2% of IBS in those patients (71.4%) who had a documented Shigella infection. The authors also found that 22.4% of patients (v 7.4% of controls) exhibited functional gastrointestinal symptoms that did not meet the Rome II criteria. These results indicate that the incidence of chronic gastrointestinal symptoms following acute gastroenteritis is similar to that reported in the UK and North America. The study by Wang and colleagues also provides the first postive association between Shigella infection and IBS, as pathogens reported in previous studies were Salmonella and Campylobacter. Taken together, these observations suggest that host, rather than microbial, factors influence the development of post-infective IBS (PI-IBS), although Campylobacter toxin may also contribute to the pathogenesis of PI-IBS. As in previous studies, the severity of the acute infection was found to be a strong risk factor for the development of PI-IBS. In contrast with a significant predisposition of females in Western studies, the study of Wang and colleagues did not identify sex as a risk factor for PI-IBS, and this is in keeping with other IBS studies from Asia. A UK study has shown that 50% of IBS patients remain symptomatic six years post-gastroenteritis and it would be of interest to determine whether there is a similar prognosis in the Beijing patients.

The study by Wang and colleagues confirmed and extended the findings from Western studies, indicating that low grade inflammation is present in patients with PI-IBS. Increased expression of interleukin (IL)-1β mRNA in rectal biopsies of PI-IBS patients confirms the recent observations of Gwee and colleagues. The authors also found increased IL-1β expression in the terminal ileum. Western studies on PI-IBS patients have been restricted to the rectosigmoid region and have not found increased mast cell numbers. However, previous studies found increased numbers of mast cells in the caecum, terminal ileum, and descending colon of IBS patients although it is not known how many of those patients had a history of gastroenteritis at the onset of IBS. Spiller et al found increased numbers of intraepithelial lymphocytes and T lymphocytes, but not mast cells, in the rectosigmoid region of PI-IBS patients and suggested that these histological findings could distinguish between IBS of infective and idiopathic aetiologies. As these studies were conducted on Western patients, it is not known whether the discrepancy in mast cell number in PI-IBS patients observed in the studies from the UK and Beijing reflect differences in patient recruitment, tissue sampling or processing, or an important biological variable related to either the infective agent or differences, for example, in microflora. Further studies are clearly necessary.

The immunohistochemical studies reported by Wang and colleagues were performed on a mixed group of IBS patients that met the Rome II criteria, consisting of patients with (48%) or without (52%) a history of gastroenteritis. They found an increased density of 5-hydroxytryptamine and substance P immunoreactive nerves surrounding mast cells in the ileum and colon of IBS patients. As there was no change in the density of calcitonin gene related peptide positive nerves, the findings unlikely reflect a generalised upregulation of nerves secondary to low grade inflammation. These data confirm the recent findings of Barbara and colleagues showing close proximity of tryptase positive mast cells and nerves along with correlation of these intimate nerve to mast cell interactions with severity of abdominal pain in IBS patients. Taken together, these two independent studies from very different regions of the globe provide strong evidence supporting a role for neuroimmune interactions in IBS. The feasibility of such interactions is provided by previous demonstrations that substance P, in low concentrations, can alter mast cell excitability and can directly influence mast cell function when released from nerves. These findings are in keeping with the demonstration that an increased rate of mast cell degranulation was found for mast cells located in close vicinity to nerves in the colonic mucosa of IBS patients. Interestingly, T lymphocytes may influence expression of neurokinin (NK)-1 receptors on mast cells and this may be important in the context of IBS where...
there is evidence of increased lymphocyte number and activation. A recent study has shown that the Th2 cytokine IL-4 upregulates NK-1 receptors on mast cells and it is therefore possible that the Th polarity of intestinal immunity could influence the nature of neuroimmune interactions in IBS. As a Th2 immune response bias exists in countries where intestinal parasitic infection is endemic, it is possible that resulting changes in neuroimmune interactions could produce differences in the pathophysiology and clinical expression of IBS between Western and Eastern societies.

There is increasing recognition of the importance of infection in the pathogenesis of IBS, and the study of Wang et al places this relationship in a broad global context. With emerging evidence supporting a role for inflammation and innate immunity activation in IBS, studies are encouraged to address the influence of the microbial environment on the epidemiology and clinical expression of IBS across the globe.


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Authors’ affiliations
S M Collins, Intestinal Diseases Research Program, Division of Gastroenterology, McMaster University Medical Centre, Hamilton, Ontario, Canada
G Barbara, Department of Internal Medicine and Gastroenterology, University of Bologna, Italy

Correspondence to: Professor S M Collins, Room 4W8, HSC, McMaster University Medical Centre, Hamilton, Ontario L8N 3Z5, Canada; scollins@mcmaster.ca

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M H Holtmann and M F Neurath

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