Cell cycle regulation in patients with intestinal metaplasia at the gastro–oesophageal junction

N J Trudgill, S K Suvarna, J A Royds, S A Riley

Background/Aims: The incidence of oesophageal adenocarcinoma is increasing rapidly and this may be related to the presence of intestinal metaplasia (IM) at the gastro–oesophageal junction (GOJ). Recent studies have distinguished two subtypes of IM at the GOJ: short segment Barrett’s oesophagus (SSBO) and IM at a normal squamo–columnar junction (IMNSCJ). Because abnormal expression of cell cycle regulators is common in cancer and precancerous states, cell cycle regulation was studied in patients with IM at the GOJ.

Methods: Biopsy samples and resected materials were identified from patients with SSBO (n = 10), IMNSCJ (n = 14), a normal SCJ with (n = 14) and without (n = 12) inflammation, conventional Barrett’s oesophagus (BO) (n = 12), and oesophageal adenocarcinoma (n = 12). Sections were stained with antibodies to p21, p27, p53, Ki67, cyclin D1, and c-erbB2 and were assessed independently by two observers, using predetermined criteria.

Results: Patients with oesophageal adenocarcinoma showed high expression of c-erbB2, p53, p27, and Ki67. Patients with BO showed expression of c-erbB2 but little expression of other markers. Greatly increased expression of cyclin D1 was seen in patients with IMNSCI. The expression of all other markers was similar in patients with IMNSCI and those with SSBO. Cyclin D1 and c-erbB2 were coexpressed in patients with SSBO and IMNSCI, and their expression was associated with the presence of p53 and p21.

Conclusions: Although the proposed aetiologies of SSBO (gastro–oesophageal reflux) and IMNSCI (Helicobacter pylori infection) differ, the cell cycle response is similar and both may have malignant potential.

METHODS

Subjects

Samples from six groups of subjects were studied. Patients with a normal SCJ were identified and subdivided into two groups: those with (n = 14) and without (n = 12) inflammation (according to the presence or absence of neutrophil polymorphs in the lamina propria). Patients with intestinal metaplasia at the gastro–oesophageal junction were identified and subdivided into two groups: those with a normal SCJ (n = 14) and those with SSBO (n = 10). A further group with conventional Barrett’s oesophagus (n = 12) was studied. Patients with dysplasia were excluded from our study. A final group comprised patients with oesophageal adenocarcinoma associated with oesophageal columnar epithelium with intestinal metaplasia (n = 12).

For the purpose of our study, a normal SCJ was defined as one that was coincident with the proximal limit of the gastric folds. Barrett’s oesophagus was defined by the combination of the endoscopic appearance of at least 3 cm of gastric-like mucosa extending above the proximal margin of the gastric folds, together with the presence of intestinal metaplasia. Those with a short segment (0.5 to 2.5 cm) of gastric-like mucosa and those with tongues of gastric-like mucosa

Abbreviations: CDK, cyclin dependent kinase; IMNSCJ, intestinal metaplasia at a normal squamo–columnar junction; PBS, phosphate buffered saline; SCJ, squamo–columnar junction; SSBO, short segment Barrett’s oesophagus
extending at least 0.5 cm above the SCJ and the presence of intestinal metaplasia were grouped together as SSBO.

**Section preparation and immunohistochemical staining**
Sections (4 μm thick) were cut and mounted on 3,3-diaminobenzidine hydrochloride as chromogen. Sections were scored, independently, by two pathologists. Sections were scored, independently, by two pathologists.

**RESULTS**

**Subjects**
Table 1 shows the demographic data of the six groups. Patients with Barrett's oesophagus, with oesophageal adenocarcinoma, and with IMNSCJ were older than patients with a normal SCJ, with or without inflammation (p < 0.05). There were no significant sex differences between the groups. As would be anticipated, patients with Barrett’s oesophagus were more likely to be taking proton pump inhibitors than those with IMNSCJ (p = 0.05). Helicobacter pylori was far more common in patients with IMNSCJ or a normal SCJ with inflammation than in patients with a normal SCJ without inflammation (p < 0.001).

**Immunohistochemistry**
Table 2 shows the expression of the six antibodies in each of the six groups.

In patients with a normal SCJ, proliferative gene expression was low. Patients with inflammation at the SCJ had increased expression of Ki67 and a modest increase in p27 and p21, but these changes were not significant. Within these combined groups, an association was found between p27 and cyclin D1 expression (p = 0.04).

Patients with conventional Barrett’s oesophagus had higher expression of c-erbB2 than those with a normal SCJ (p = 0.04). Surprisingly, the expression of proliferative markers was low.

However, patients with oesophageal adenocarcinoma showed high c-erbB2 expression, together with increased p53 expression, compared with patients with a normal SCJ (p < 0.01), SSBO/IMNSCJ (p = 0.02), or Barrett’s oesophagus (p < 0.01). p21 expression appeared to be increased, although the results were not significant. Patients with adenocarcinoma also showed higher expression of p27 than did those with a normal SCJ (p = 0.02) or Barrett’s oesophagus (p < 0.01), and increased Ki67 expression compared with patients with a normal SCJ (p = 0.02), SSBO/IMNSCJ (p < 0.01), or Barrett’s oesophagus (p = 0.03).

In patients with IMNSCJ and SSBO, c-erbB2 expression was similar to that seen in patients with Barrett’s oesophagus, but p27 expression was significantly increased (combined data from patients with SSBO and IMSCJ compared with patients with Barrett’s oesophagus; p = 0.02). When the two groups were compared, greatly increased expression of cyclin D1 was found in patients with IMNSCJ than in those with SSBO (p = 0.05). However, there were no other significant differences in expression scores between these two groups.

In the combined SSBO and IMNSCJ group, cyclin D1 and c-erbB2 expression appeared to be associated. Seventy nine per cent of sections that were cyclin D1 positive were also c-erbB2 positive, whereas 22% of cyclin D1 negative sections were c-erbB2 positive (p = 0.03). Combined cyclin D1 and c-erbB2 expression paralleled the expression of both p53 and p21. Sixty per cent of cyclin D1 positive sections were p53 positive, whereas only 11% of cyclin D1 negative sections were p53 positive (p = 0.05). Sixty nine per cent of c-erbB2 positive sections were also p21 positive, but only 20% of c-erbB2 negative sections were p21 positive (p = 0.05).

**DISCUSSION**
Barrett’s oesophagus is the principal risk factor for the development of oesophageal adenocarcinoma. However, adenocarcinoma may also arise in short segments of gastric-like mucosa with intestinal metaplasia that fail to reach conventional criteria for Barrett’s oesophagus. Recent studies have distinguished two subtypes of intestinal metaplasia at the gastro-oesophageal junction: SSBO and IMNSCJ. Preliminary natural history data confirm that dysplasia and carcinoma can develop in such segments, but the risk of malignant transformation in comparison with patients with conventional Barrett’s oesophagus has not been assessed.

In other parts of the gastrointestinal tract, evidence of stepwise oncogene activation with loss of tumour suppressor...
gene function is seen during the progression from normal, through dysplastic states, to frank neoplasia. Therefore, our study chose to consider cell cycle protein expression in the groups described above by means of immunohistochemistry in an attempt to clarify whether SSBO, IMNSCJ, and Barrett’s oesophagus have a similar pathophysiology and malignant potential.

The markers used in our study examine several different aspects of the cell cycle. p53 is expressed in response to DNA damage and regulates cell division by preventing progression from G1 to S phase, allowing DNA repair to take place. p21 is a cyclin dependent kinase (CDK) inhibitor induced by wild-type p53 to downregulate cell division. p27 is also a CDK inhibitor and regulates the cell cycle, but its expression is not dependent on p53. Cyclin D1, in collaboration with CDKs, facilitates progression through the G1 phase of the cell cycle and may act as an oncogene. Ki67 is a nuclear antigen that is homologous to the epidermal growth factor receptor and is regarded as a protooncogene, increasing cellular proliferation.

The presence of inflammation was associated with an increase in Ki67, indicating increased cellular proliferation and a modest increase in cell cycle regulators: p53, p21, and p27.

Previous publications have revealed abnormalities of several markers of the cell cycle in relation to progression from intestinal metaplasia to dysplasia and adenocarcinoma in patients with Barrett’s oesophagus. Mutations of p53 have been reported in Barrett’s oesophagus with intestinal metaplasia alone or with low grade dysplasia, suggesting that this may be a relatively early event in the progression to adenocarcinoma. Increased p21 expression is also found in Barrett’s oesophagus scored indefinite for dysplasia or low grade dysplasia compared with intestinal metaplasia alone, again suggesting that this is an early event. p27 expression increases in dysplastic Barrett’s oesophagus compared with metaplasia, but it is inactivated in adenocarcinoma, and its loss is associated with a poor outcome. Cyclin D1 expression is also common in Barrett’s oesophagus without dysplasia, and is associated with a higher risk of malignant progression. Finally, overexpression of the protooncogene c-erbB2 may be an early event because it has been described in intestinal metaplasia, dysplasia, and adenocarcinoma.

In our present study, we have examined cell cycle markers in tissue samples from patients with intestinal metaplasia in the lower oesophagus and a range of controls. As anticipated, little proliferative activity was seen in patients with a normal SCJ. The presence of inflammation was associated with an increase in Ki67, indicating increased cellular proliferation and a modest increase in cell cycle regulators: p53, p21, and p27. Within these groups, an association was seen between p27 and cyclin D1. This is perhaps not surprising because p27 controls cyclin D1 activity by complexing with it, thereby inhibiting cell cycle associated kinase activity.

In patients with Barrett’s oesophagus, increased expression of c-erbB2 was apparent without an increase in proliferative markers. This unexpected finding can be explained in part by the fact that most of the patients with Barrett’s oesophagus were taking proton pump inhibitors at the time of their endoscopy. Intermittent exposure to acid has been shown to promote proliferation and reduce differentiation in Barrett’s oesophagus cell cultures, and effective acid suppression, judged by 24 hour pH monitoring, has been shown to reduce proliferative activity in patients with Barrett’s oesophagus.

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<th>Table 2 Marker expression in the six study groups</th>
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Mean score (range; percentage + or more).
BO, Barrett’s oesophagus; IMNSCJ, intestinal metaplasia at a normal squamo–columnar junction; OAC, oesophageal adenocarcinoma; SCJ, squamo–columnar junction; SSBO, short segment Barrett’s oesophagus.
In patients with adenocarcinoma, high c-erbB2 expression was associated with increased proliferative markers and loss of cell cycle control, and these results broadly parallel previously published data.15 17 26

With the exception of cyclin D1, marker expression in patients with SSBO and IMNSCJ was similar, and the expression of c-erbB2 was similar to that seen in patients with Barrett’s oesophagus. However, in patients with IMNSCJ (H pylori associated intestinal metaplasia), there was a pronounced increase in expression of the potential oncogene, cyclin D1, and this appeared to be associated with upregulation of its inhibitor, p27. Others have also suggested that H pylori may activate cyclin D1.30 In patients with SSBO, although cyclin D1 expression was not increased, p27 appeared to be upregulated. Furthermore, within both groups, cyclin D1 and c-erbB2 were associated and paralleled the expression of both p53 and p21. It is possible that external factors are stimulating metastatic cells to proliferate and mutations to genes controlling the cell cycle, such as p53, may subsequently result in the development of dysplasia and adenocarcinoma.

We appreciate that our study has limitations. First, we recognise that our patient classifications are based on the histological finding of intestinal metaplasia in relatively small biopsy samples. We realised that metaplasia can be patchy and recognise that our patient classifications are based on the histological finding of intestinal metaplasia in relatively small biopsy samples. We realise that metaplasia can be patchy and minimise the differences between the two groups. Second, the numbers studied were relatively small, limiting the power of our conclusions. Further studies are clearly indicated.

In conclusion, the results of our present study have shown that oesophageal adenocarcinoma is associated with deregulation of the cell cycle and apoptotic control. With the exception of cyclin D1, expression of the markers studied in IMNSCJ and SSBO was similar, and c-erbB2 expression was comparable to that seen in Barrett’s oesophagus. An increase in cyclin D1 expression in apparent in patients with IMNSCJ, and probably relates to the associated H pylori infection. We feel that although the proposed aetiologies of SSBO (gastric-oesophageal reflux) and IMNSCJ (H pylori infection) differ, the cell cycle response appears to be similar, and both may have malignant potential.

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Authors’ affiliations

N J Trudgill, S A Riley, Department of Gastroenterology, Northern General Hospital, Sheffield SS 7AU, UK
S K Suvarna, Department of Histopathology, Northern General Hospital
J A Royds, University of Otago, New Zealand

REFERENCES


The first study to test for DNA adducts as a marker for intrahepatic cancer of the bile duct has lent more support to the hypothesis that toxins damaging DNA lead to cancer. Greater exposure to environmental toxins may explain the rapid rise in deaths from this cancer in Western countries over the past 30 years, suggest the authors.

Patients with the cancer had more DNA adducts in samples of their tumour or adjacent to their tumour than patients without (median relative amounts/10^8 nucleotides: tumour DNA 7.2 (range 1.8–48.4); tumour adjacent DNA 8.6 (1.2–51.6) v non-cancer DNA 2.9 (0.6–11.5)). The pattern and density of DNA adducts differed between tumour DNA and tumour adjacent DNA and between each of these and non-cancer DNA.

Tumour tissue was taken from 32 men with primary intrahepatic cholangiocarcinoma and tumour adjacent tissue from 28 of them. Cystic ducts were obtained from seven cancer free patients during surgery for gall stones. Adducts were detected in DNA extracted from each sample by 32P postlabelling assays, TLC separation, and autoradiography. Positive and negative controls were included.

DNA adducts are modified DNA bases formed when electrophilic carcinogens or their metabolites bind to DNA. They are key to chemically induced mutagenesis through misrepair or absence of repair and are markers for chemical mutagenesis, but not hitherto in bile duct tissue, even though the liver and biliary system metabolise carcinogens. Intrahepatic bile duct cancer is the second commonest liver cancer worldwide, but the cause of its prominence is unknown.

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