Oncogenes and onco-suppressor gene in adenocarcinoma of the oesophagus

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Abstract
While the activation of the proto-oncogenes has been implicated in the development and progression of cancer of many tissues, the role of oncogenes in the development of oesophageal adenocarcinoma has not been defined. Fifteen patients who had undergone resection for oesophageal adenocarcinoma and 15 who had undergone oesophagectomy or biopsy for Barrett’s oesophagus were studied. The latter patients also had adjacent normal gastric mucosa biopsied for comparison with the metaplastic oesophageal mucosa. The mucosal samples were snap frozen and subsequently stained with monoclonal antibodies to the following oncogene associated proteins; c-erbB2 (neu and CE-1) (external domain), c-erbB2 (NCL-CB11) (internal domain), c-src, c-ras, c-myc, c-fos, c-jun, and the onco-suppressor gene – p53. All tumours were well or moderately differentiated adenocarcinomas arising from the lower third of the oesophagus. Eleven specimens showed strong membranous staining with both c-erbB2 (neu) and c-erbB2 (CBL-CB11). Seven specimens showed strong nuclear staining with p53 onco-suppressor gene. Three specimens were positive for c-ras and c-src, and two were positive for c-jun. In Barrett’s epithelium, nine specimens were positive for c-erbB2 (neu and CB11), three were positive for c-src, two were positive for c-ras and c-jun, and one was positive for c-fos. Two of the gastric mucosal biopsy specimens expressed c-erbB2 weakly but no other oncogenes were found. The frequency of positive staining for c-erbB2 is very high, compared with the expression of these genes in other tumours. It is also concluded that errors in the onco-suppressor gene p53, and especially in the external and internal domains of c-erbB2, which is also often expressed in Barrett’s mucosa, may be implicated in the development of adenocarcinoma of the oesophagus.

In the past decade considerable progress has been made in the identification of genes that encode proteins controlling the complex cascade of processes regulating the proliferation and differentiation of cells. The proliferation of normal and malignant cells is regulated by both inhibitory and stimulatory control molecules, derived from tumour suppressor genes and proto-oncogenes respectively. A tendency towards cancer-associated loss of normal growth control can result from loss or inactivation of both copies of a tumour suppressor gene or from mutation, with amplification of hyperactivation of one or both of the two copies of one, or more, proto-oncogenes.

In addition, recent reports have suggested that the degree and type of oncogene expression in some tumours (breast, ovary, pancreatic, stomach, and colon) may have prognostic importance.

Oncogenes can be divided into three categories according to their subcellular localisation and their function in cellular transformation – growth factor oncogenes; signal transducer genes; and oncogenes encoding nuclear proteins. We, and others, have previously reported abnormal expression of growth factors and their receptors in oesophageal adenocarcinomas. It therefore seemed pertinent to assess also the expression of membrane associated signal transducer genes and ‘nuclear oncogenes’, particularly since abnormal expression of these oncogenes may be implicated in the development of Barrett’s oesophagus (a precursor of adenocarcinoma) as well as adenocarcinomas.

Patients and methods
Fifteen patients who had undergone oesophageal resection for oesophageal adenocarcinoma (mean age 67 years, range 51–76) were assessed in addition to 15 others who had undergone biopsy or resection for Barrett’s oesophagus (mean age 64 years, range 43–80).

Forty five specimens were evaluated, comprising 15 resected adenocarcinomas, 15 cases of Barrett’s oesophagus, and paired biopsy specimens of normal gastric mucosa from the latter 15 patients.

Specimens were snap frozen and subsequently cut to 6 μm sections. Routine histological assessment was performed using haematoxylin and eosin stained sections. The degree of differentiation of the cancers was classified according to previous criteria. The lesions of Barrett’s oesophagus were graded for metaplasia and dysplasia according to the criteria of Whitehead and Riddell respectively.

The following monoclonal antibodies were used to assess expression of appropriate genes: internal domain of c-erbB2, NCL-CB11 (IgG, Novocastra); external domain of c-erbB2 neu TA1 (IgG, Du Pont); anti-pan ras p21 (IgG, Du Pont); anti-p53 (IgG, gift from Professor D Lane, Biochemistry Dept, Dundee); c-src (IgG, Cambridge Research Biochemicals); c-jun (IgG, Cambridge Research Biochemicals Ltd); c-fos (IgM Bionuclear Services Ltd); and c-myc (IgG, Cambridge Research Biochemicals).

Sections were fixed in acetone for five minutes at room temperature, and then stained according to the indirect immunoperoxidase technique with biotinylated mouse IgG (DAKO) as a...
secondary antibody followed by streptavidine, and this product was visualised by diaminobenzidine as described previously.5

Positive control sections included lymph node containing lymphoma (c-myc, c-jun, and c-fos demonstrated staining confined to the nucleus of lymphocytes and stromal cells), mammary and gastric adenocarcinomas (c-erbB2 internal and external domains and c-src demonstrated strong membranous staining in epithelial tissues) (c-ras demonstrated moderate cytoplasmic staining in epithelial tissues), colonic adenocarcinomas (p53 demonstrated strong nuclear staining in epithelial cells).

Negative control sections for each oncogene were processed immunohistochemically without the primary antibody.

Immunohistochemical staining of cell membranes (C-erbB2, C-src, cytoplasmic (c-ras), or nucleus (c-myc, c-fos, c-jun and p53) was scored by assessing both the intensity of staining and the proportion of cells stained, based on a modification of the technique of Corbett et al.21

Cytoplasmic staining was not considered to be positive for the purposes of semi-quantitation of c-erbB2. When cytoplasmic c-erbB2 staining was present it was noted separately (Table I).

Briefly, intensity scored on a four point scale: no staining=0; weak staining=+; moderate staining=++; and intense staining=+++.

The proportion of cells stained was also scored on a four point scale: no cells staining=0; 1–39% staining =I; 40–69% staining =II; 70–100% staining =III.

Two individuals assessed each section and there was complete agreement in 95% of cases with regard to the intensity and pattern of staining. In the few sections in which scores differed, the two observers agreed a common score.

### Table I: Staining patterns of monoclonal antibodies to oncogene and anti-oncogene associated epitopes

<table>
<thead>
<tr>
<th>Oncogene</th>
<th>Dilution</th>
<th>Tissue stained</th>
<th>Staining pattern</th>
</tr>
</thead>
<tbody>
<tr>
<td>NEU</td>
<td>1/10</td>
<td>Epithelium</td>
<td>Membranous and cytoplasmic</td>
</tr>
<tr>
<td>CB11</td>
<td>1/5</td>
<td>Epithelium</td>
<td>Membranous and cytoplasmic</td>
</tr>
<tr>
<td>c-src</td>
<td>1/100</td>
<td>Epithelium</td>
<td>Cytoplasmic and membranous</td>
</tr>
<tr>
<td>c-ras</td>
<td>1/20</td>
<td>Epithelium</td>
<td>Cytoplasmic</td>
</tr>
<tr>
<td>c-jun</td>
<td>1/50</td>
<td>Epithelium</td>
<td>Cytoplasmic and nuclear</td>
</tr>
<tr>
<td>c-fos</td>
<td>1/10</td>
<td>Epithelium and stromal cells</td>
<td>Nuclear</td>
</tr>
<tr>
<td>p53</td>
<td>1/10</td>
<td>Epithelium</td>
<td>Cytoplasmic and membranous</td>
</tr>
</tbody>
</table>

### Table II: Percentage of positive expression of oncogene and anti-oncogene associated antigens in the oesophagus in health and disease

<table>
<thead>
<tr>
<th>Oncogene</th>
<th>Adenocarcinoma (%)</th>
<th>Barrett’s mucosa (%)</th>
<th>Gastric mucosa (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Neu</td>
<td>(+ + +) 73</td>
<td>(+) 60</td>
<td>(+) 13</td>
</tr>
<tr>
<td>CB11</td>
<td>(+ + ) 73</td>
<td>(+) 60</td>
<td>(+) 7</td>
</tr>
<tr>
<td>c-src</td>
<td>(+ + ) 20</td>
<td>(+) 20</td>
<td>(+) 7</td>
</tr>
<tr>
<td>c-ras</td>
<td>(+ + ) 20</td>
<td>(+) 13</td>
<td>0</td>
</tr>
<tr>
<td>c-jun</td>
<td>(+ + ) 20</td>
<td>(+) 13</td>
<td>0</td>
</tr>
<tr>
<td>c-fos</td>
<td>(+ + ) 7</td>
<td>(+) 7</td>
<td>0</td>
</tr>
<tr>
<td>c-myc</td>
<td>(+ + ) 0</td>
<td>(+ + ) 0</td>
<td>0</td>
</tr>
<tr>
<td>p53</td>
<td>(+ + +) 46</td>
<td>[II] 0</td>
<td>0</td>
</tr>
</tbody>
</table>

*Average staining intensity in positive sections in brackets; (+) = weak, (+ + ) = moderate, (+ + +) = intense.
†Average staining intensity in positive sections in square brackets; [I] = 1–39%, [II] = 40–69%, [III] = 70–100%

### Results

All adenocarcinomas were well or moderately differentiated. Of the 15 cases of Barrett’s oesophagus, eight contained mainly (>90% mucosa) intestinal type mucosa, three contained mainly (>90%) gastric type mucosa, and four contained both types of epithelium. All gastric mucosal biopsy specimens were histologically unremarkable.

The pattern of positive expression of different oncogenes was variable. For example, c-erbB2 (neu and CB11) showed both membranous distribution and minimal cytoplasmic staining in most positive cells. C-erbB2 cytoplasmic staining was not included in the assessment of staining density; this included only membranous staining as suggested previously.21 C-src showed predominantly cytoplasmic distribution with minimal membranous staining. Similarly, c-ras and c-jun had a cytoplasmic pattern of reactivity whereas c-fos and p53 were both nuclear (Table I). C-myc failed to react positively with any of the sections under study, although a control section of lymph node containing lymphoma showed positive staining (Table II).

The density of staining for all oncogenes was variable throughout the sections (Table II). The staining pattern also varied from one cancer to another and one section of Barrett’s oesophagus to another.

The proportion of cells positively stained also varied from one section to another, but to a lesser degree.

### Background Staining

Background staining was minimal or absent in most sections and did not therefore interfere with semiquantitation. The following antibodies achieved particularly specific staining with an absence of background staining: neu, CB11, p53, c-ras, and c-fos. The following antibodies had some diffuse background staining usually localised in superficial epithelial cells or stromal tissue; c-jun, c-myc, and c-src. With empirical methods of dilution and application of fetal calf serum (1/25), before incubation with the primary antibodies we were able to minimise the background staining.

### Membrane Bound Oncogenes

Both the internal and external domain of c-erbB2 (Fig 1) were coherently expressed in most sections from oesophageal carcinomas and Barrett’s oesophagus (73% and 60%, respectively).

Seven sections with Barrett’s intestinal type mucosa stained positively. In addition to the membranous staining, there was mild-moderate cytoplasmic staining in 6 of 12 sections of positively staining carcinomas and 5 of 9 sections of Barrett’s oesophagus (one with mild dysplasia). There was no cytoplasmic staining in normal mucosa. In two sections, however, weak membrane staining was present in superficial and foveolar type cells of the gastric crypts.

### Cytoplasmic Oncogenes

C-ras (Fig 2) was expressed in 20% of oesophageal

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healthy gastric mucosa also expressed c-src weakly in the cytoplasm of epithelial cells.

NUCLEAR ONCOGENES
C-myc was not demonstrated in any oesophageal sections. C-fos (Fig 4) was found in one case of Barrett’s oesophagus (gastric type metaplasia). C-jun (Fig 5) was found in 20% of oesophageal cancers and in 13% of Barrett’s oesophagus. The pattern of expression was mainly cytoplasmic rather than nuclear.

P53 ANTI-ONCOGENE
Seven of 15 (Fig 6) oesophageal adenocarcinomas stained strongly positive in the nuclei with p53 antiserum. All staining was confined to the malignant mucosa with no staining of the stromal nuclei. In the sections that stained positively, approximately 40–70% of malignant cells were stained with p53. One section of non-dysplastic Barrett’s mucosa had occasional faint nuclear staining.

Discussion
The results of the present study show that c-erbB2 internal and external domains are coherently expressed in oesophageal adenocarcinomas and Barrett’s oesophagus. The coherent expression of internal and external domains of c-erbB2 has also been reported in mammary carcinoma.11 The erbB2 protein on cell membranes has a strong correlation with its corresponding mRNA which, in turn, is produced by amplification of the erbB2 proto-oncogene.11,12

The present study has shown a very high incidence of membranous erbB2 expression in both malignant and Barrett’s oesophageal tissue (73% expression and 60%, respectively) compared with the lower frequency of expression in other gastrointestinal cancers (gastric 30% and colonic 50%).13,17,27,30 This finding is interesting because we have recently reported an increased expression of epidermal growth factor receptors (EGF-R), which also have tyrosine kinase activity, in both oesophageal cancer and Barrett’s epithelium.11 However, the COOH domain of EGF-R and c-erbB2 exert different transforming activity.15 The erbB2 COOH domain has a positive stimulatory effect on erbB2 oncogenic activity, perhaps accounting for the fact that tumours expressing erbB2 have both a poorer short term (<5 years) as well as long term (>10 years) prognosis.11,15,17,26,28,30 Moreover, tumours with erbB2 expression have been reported to metastasise readily.11,30 In this context, we have shown a paucity of c-erbB2 expression in epithelial membranes in normal gastric mucosa, with greater levels found in Barrett’s oesophagus and the greatest levels of c-erbB2 expression on oesophageal carcinomas. In this respect membranous c-erbB2 expression is a marker for malignant potential, however, it has been suggested that cytoplasmic c-erbB2 expression, noted also in six sections of adenocarcinoma, may have less prognostic importance than membranous expression.11 Whether this staining

carcinomas and 14% of Barrett’s oesophagus (both of the latter had intestinal metaplasia). C-src (Fig 3) was expressed in 20% of oesophageal carcinomas and also of Barrett’s oesophagus (2 of 3 patients had intestinal metaplasia). One case of

Figure 1: (A) C-erbB2 neu (external domain) in adenocarcinoma of the oesophagus (original magnification ×450). (B) C-erbB2 CB-11 (internal domain) in adenocarcinoma of the oesophagus (original magnification ×450). (C) C-erbB2 neu in Barrett’s gastric type mucosa (original magnification ×450).
represents extracellular domain fragment secreted by native c-erbB gene is uncertain.11

There are many possible methods by which wild type p53 could result in growth suppression,12 however, one recently reported mechanism is by selective down regulation of expression of the proliferating cell nuclear antigen.12 Therefore wild type p53 seems to exert essential inhibitory effects on cell proliferation, and therefore mutation of p53 may result in deregulation of cellular proliferation. Mutant p53 was shown in a similar proportion of sections of oesophageal adenocarcinoma (46%) compared with previous reports of other alimentary tumours (colonic carcinoma 35%).44,45 We were unable to demonstrate unequivocal p53 staining in (premalignant) Barrett’s mucosa, irrespective of the presence or absence of dysplasia.46 However, unlike erbB2, expression of which develops early in the neoplastic process, p53 mutation is usually a late step in carcinogenesis.47 It is possible that the failure of this regulatory ‘anti-oncogene’ results in unopposed growth stimulation by the EGF-R and erbB2 receptor which is ‘normally limited’ in the nucleus by p53.46

We found c-ras in 20% of oesophageal carcinomas and 14% of Barrett’s oesophagus. These values are lower than the frequency of expression in other carcinomas (colonic cancer 25%, pancreatic and breast cancer 30%, and gastric cancer 35%).48,49 The relatively low frequency in oesophageal carcinoma may, in part, explain the undetectable levels of ras mRNA reported previously.49

C-src was expressed in 20% of oesophageal cancers and Barrett’s oesophagus, compared with 30% of gastric adenocarcinomas reported previously.44 We have also shown that normal gastric mucosa may also express c-src, as reported previously.44 The protein is predominantly expressed in the Barrett’s epithelial cell cytoplasm but also to a lesser extent in the cell membranes. C-src is an oncogene which encodes a cytoplasmic protein with tyrosine kinase activity and may influence the regulation of growth control by affecting the expression of specific genes for epidermal growth factor receptors.50 In addition c-src may deregulate cell adhesion and anchorage dependent growth control, thereby maintaining cells in a proliferative state.44,45

C-myc was absent in all sections of the oesophageal adenocarcinomas. In this context it has been reported that c-myc has a short half life,44 furthermore c-myc expression is indirectly related to lack of tissue differentiation51 and may not have been optimally expressed in our carcinomas.

Only one case of Barrett’s gastric type mucosa was positive for c-fos, perhaps because c-fos is very unstable and has strong negative regulation.52

C-jun was expressed most frequently of all the nuclear oncogenes. Unlike other nuclear genes, c-jun undergoes positive autoregulation. It has been reported that the degree of c-jun expression is involved in converting short term, proliferation inducing signals into long term memory, thus controlling the number of cell divisions.53 The cytoplasmic expression of c-jun in the present
Figure 5: Expression of c-jun in Barrett's gastric type mucosa (weak staining of the apical cytoplasm is visible) (original magnification ×700).

Figure 6: Expression of p53 oncogene in the nucleus of oesophageal adenocarcinoma (original magnification ×450).

Oncogenes in oesophageal adenocarcinoma

study may represent deregulation of the c-jun oncogene in oesophageal diseases as has been shown with other nuclear oncogenes like c-myc. It has been suggested that expression of one oncogene alone may not result in frank malignancy because cancers, as a rule, result from the chance occurrence in one cell of several independent genetic changes. It is likely that the synchronous expression of oncogenes may act synergistically, gradually overcoming the negative feedback regulation of mitogenesis. In this respect, the expression of an abnormal p53 anti-oncogene, if it occurs at all, may represent a late step in the development of oesophageal cancer, although mutant p53 has occasionally been found in 'normal' epithelia.

In conclusion, we have shown that a variety of oncogenes are expressed in oesophageal adenocarcinoma and Barrett's oesophagus. The frequency of expression of c-erbB2 and p53 suggests that the oncogene and an abnormal anti-oncogene may also be important in the development of oesophageal adenocarcinoma. We are presently assessing the prognostic potential of these oncogenes, especially since the incidence of adenocarcinoma is increasing in Tayside.


