Classification and localisation of carcinoembryonic antigen (CEA) related antigen expression in normal oesophageal squamous mucosa and squamous carcinoma

D S A Sanders, C A Wilson, F J Bryant, J Hopkins, G D Johnson, D M Milne, M A Kerr

Abstract
Using a panel of carcinoembryonic antigen (CEA) related antibodies in normal oesophageal squamous mucosa CEA expression is present on suprabasal squames localised to the cell membrane. Immunoblotting shows that this positivity is predominantly due to a glycoprotein of around 180 kDa representing CEA itself. Positivity in squamous carcinomas is confined to cells in foci of squamous differentiation. A shift from membranous localisation to predominant cytoplasmic overexpression is shown between normal and malignant squames using confocal microscopy. The recognition of an adhesive role for CEA and a role in enhancing distant metastases in those tumours expressing CEA highlights the importance of recording CEA expression and changes in subcellular distribution between normal and malignant tissues; CEA expression in oesophageal squamous mucosa has not been well recognised previously and changes in expression may prove of great significance in the spread and dissemination of squamous carcinoma.

Recent advances in molecular biology and gene cloning have resulted in great improvements in the understanding of the carcinoembryonic antigen (CEA) related family of glycoproteins leading to a reassessment of the clinical outlook for CEA and giving insight into the function of these molecules. The CEA gene family is now known to belong to the immunoglobulin superfamily and in particular the C-2 set of adhesion molecules; cell adhesion properties are well established for CEA and its classic related cross reacting antigens, non-specific cross reacting antigen (NCA), and biliary glycoprotein. Binding may be homophilic (CEA to CEA) or heterophilic (NCA to CEA), is neither calcium or temperature dependent, and is capacitated by membranous expression on cells.

In the gastrointestinal tract most attention has focused on CEA expression in the normal colonic and gastric mucosa and adenocarcinomas. CEA expression in squamous mucosa and squamous carcinoma is not widely recognised and has received little attention. We recently reported expression of a CEA related antigen on the cell surface of squames in normal cervical mucosa, which suggests a key role in maintaining the integrity of the mucosa through an adhesive function. Any change in the pattern of expression and subcellular distribution of CEA, as has been reported between normal colonic mucosa and adenocarcinoma, may be of significance in tumour dissemination.

We therefore have now extended our previous study to classify and localise CEA related antigen positivity in normal...
Methods

IMMUNOHISTOCHEMISTRY

Sections cut from archive paraffin wax blocks from 30 cases of invasive oesophageal squamous carcinoma and from 20 blocks of endoscopic biopsy specimens of normal oesophageal mucosa from non-cancer patients were pretreated with 0.1% trypsin in TRIS buffered saline at pH 7.6 at room temperature for 25 minutes and stained with a panel of anti-CEA antibodies — polyclonal anti-CEA (DAKO) at 1/200 dilution; monoclonal anti-CEA A5B7 (DAKO) at 1/50 dilution; monoclonal anti-CEA IM10 6:2 (Scottish Antibody Production Unit) 1/50 dilution.

All sections were developed with the appropriate second antibody and an ABC Vectastain kit (Vector Laboratories, Peterborough). Control sections were prepared by missing out the anti-CEA.

IMMUNOBLOTTING

Fresh normal mid-oesophageal endoscopic biopsy specimens were used from two patients. Tissue was finely chopped re-suspended in phosphate buffered saline at a concentration of 100 mg wet tissue/ml and then homogenised using a Polytron homogeniser at half speed for two minutes. Triton X-100 was added to a final concentration of 1% vol/vol and the extract rotated for one hour at 4°C. After centrifugation the supernatent was removed for further study. Proteins were separated by SDS-polyacrylamide gel electrophoresis on 10% polyacrylamide gels by the method of Laemmli and stained with polyclonal anti-CEA. Extracts of neutrophil membrane were run in adjacent lanes as a positive control (prepared as in reference 7).
CONFOCAL MICROSCOPY
Six µm sections were cut from paraffin wax blocks of four cases of normal squamous mucosa and four squamous carcinomas. Sections were incubated in methanol in 0-3% hydrogen peroxide at room temperature, washed, and trypsinised. Sections were incubated with 10% normal goat serum, washed, and incubated with polyclonal anti-CEA at 1:200 dilution. After further washing the fluorescent conjugate was added 1:40 in phosphate buffered saline. Nuclei were counterstained with propidium iodide and sections mounted in Glycergel (DAKO). Control sections were prepared by missing out the anti-CEA. Sections were viewed on a Bio-Rad 500 Laser Scanning Confocal Microscope System attached to a Leitz SM-Lux fluorescence microscope.

Results
IMMUNOHISTOCHEMISTRY
In normal squamous mucosa CEA related expression was seen on the cell membrane of a broad zone of suprabasal squames with both the cytoplasm and in contrast with CEA expression in normal colon, which is on the luminal surface of cells and therefore has little part to play in maintaining mucosal integrity.4 Bacterial colonisation in the gut is known to be influenced, however, by bacterial ligand-CEA interactions.11 Expression of CEA on suprabasal (mature) squames and restricted to cells in squamous pearls in oesophageal carcinoma confirms that, as in the cervix, CEA expression, in particular membranous expression, is a marker of squamous differentiation.12 Using confocal microscopy we have shown that there is a change from membranous to predominantly cytoplasmic overexpression between normal and neoplastic squames. The differences in tissue specific CEA expression can largely be explained by the finding of widely varying levels of transcription activity within the promoter region of the CEA genes, although control of this element is poorly understood.13 Anchorage of CEA into the cell membrane is through replacement of the hydrophobic M domain on CEA with a glycosyl phosphatidylinositol moiety. Membrane bound CEA can be released through the action of specific phospholipases and it is possible that these two mechanisms interact to determine the amount of CEA expressed on cells or released into the serum, or both.14 Raised serum concentrations of CEA have been noted in squamous carcinomas.15 16 Membranous upregulation of CEA is recognised in colonic adenocarcinoma and hence CEA may have a role in promoting the metastasis of adenocarcinoma. In squamous carcinoma loss of CEA expression and a shift to cytoplasmic expression may adversely influence cellular cohesion; CEA could only have a putative influence in the metastasis of squamous carcinoma in tumours where a significant proportion of cells show membranous expression. The absence of CEA related antigen in oesophageal mucosa is a glycoprotein of around 180 kDa known to represent a true CEA. A broad band is seen on immunoblotting because the polyclonal anti-CEA recognises CEA with subtle differences of glycosylation and hence molecular weight; the core peptide of CEA itself is only 80 kDa, carbohydrate structures accounting for up to 60% of the mass of the mature glycoprotein. Variable glycosylation is also responsible for much of the complex immunogenicity of the CEA superfamily of molecules. CEA of 180 kDa is also expressed in normal cervical squamous mucosa but is slightly higher than that expressed by standard adenocarcinoma (165 kDa).3 Immunoblotting of adenocarcinoma shows additional bands from 55–120 kDa and in cervical squamous mucosa 18-5-66 kDa glycoproteins have been reported representing various NCAs.3 8 CEA 180 acts as an accessory adhesion molecule mediating interactions between tumour cells and endothelial basement membrane9 and CEA interacts with liver Kupffer cells promoting distant metastasis.10 Membranous expression of CEA in the normal oesophagus suggests a key adhesive part in maintaining the integrity of the mucosa. This is in contrast with CEA expression in normal colon, which is on the luminal surface of cells and therefore has little part to play in maintaining mucosal integrity.4 Bacterial colonisation in the gut is known to be influenced, however, by bacterial ligand-CEA interactions.11 Expression of CEA on suprabasal (mature) squames and restricted to cells in squamous pearls in oesophageal carcinoma confirms that, as in the cervix, CEA expression, in particular membranous expression, is a marker of squamous differentiation.12 Using confocal microscopy we have shown that there is a change from membranous to predominantly cytoplasmic overexpression between normal and neoplastic squames. The differences in tissue specific CEA expression can largely be explained by the finding of widely varying levels of transcription activity within the promoter region of the CEA genes, although control of this element is poorly understood.13 Anchorage of CEA into the cell membrane is through replacement of the hydrophobic M domain on CEA with a glycosyl phosphatidylinositol moiety. Membrane bound CEA can be released through the action of specific phospholipases and it is possible that these two mechanisms interact to determine the amount of CEA expressed on cells or released into the serum, or both.14 Raised serum concentrations of CEA have been noted in squamous carcinomas.15 16 Membranous upregulation of CEA is recognised in colonic adenocarcinoma and hence CEA may have a role in promoting the metastasis of adenocarcinoma. In squamous carcinoma loss of CEA expression and a shift to cytoplasmic expression may adversely influence cellular cohesion; CEA could only have a putative influence in the metastasis of squamous carcinoma in tumours where a significant proportion of cells show membranous expression. The absence of CEA related antigen in oesophageal mucosa is a glycoprotein of around 180 kDa known to represent a true CEA. A broad band is seen on immunoblotting because the polyclonal anti-CEA recognises CEA with subtle differences of glycosylation and hence molecular weight; the core peptide of CEA itself is only 80 kDa, carbohydrate structures accounting for up to 60% of the mass of the mature glycoprotein. Variable glycosylation is also responsible for much of the complex immunogenicity of the CEA superfamily of molecules. CEA of 180 kDa is also expressed in normal cervical squamous mucosa but is slightly higher than that expressed by standard adenocarcinoma (165 kDa).3 Immunoblotting of adenocarcinoma shows additional bands from 55–120 kDa and in cervical squamous mucosa 18-5-66 kDa glycoproteins have been reported representing various NCAs.3 8 CEA 180 acts as an accessory adhesion molecule mediating interactions between tumour cells and endothelial basement membrane9 and CEA interacts with liver Kupffer cells promoting distant metastasis.10 Membranous expression of CEA in the normal oesophagus suggests a key adhesive part in maintaining the integrity of the mucosa. This is in contrast with CEA expression in normal colon, which is on the luminal surface of cells and therefore has little part to play in maintaining mucosal integrity.4 Bacterial colonisation in the gut is known to be influenced, however, by bacterial ligand-CEA interactions.11 Expression of CEA on suprabasal (mature) squames and restricted to cells in squamous pearls in oesophageal carcinoma confirms that, as in the cervix, CEA expression, in particular membranous expression, is a marker of squamous differentiation.12 Using confocal microscopy we have shown that there is a change from membranous to predominantly cytoplasmic overexpression between normal and neoplastic squames. The differences in tissue specific CEA expression can largely be explained by the finding of widely varying levels of transcription activity within the promoter region of the CEA genes, although control of this element is poorly understood.13 Anchorage of CEA into the cell membrane is through replacement of the hydrophobic M domain on CEA with a glycosyl phosphatidylinositol moiety. Membrane bound CEA can be released through the action of specific phospholipases and it is possible that these two mechanisms interact to determine the amount of CEA expressed on cells or released into the serum, or both.14 Raised serum concentrations of CEA have been noted in squamous carcinomas.15 16
CEA related antigen expression in oesophageal squamous mucosa and squamous carcinoma

from cells at the periphery of differentiated squamous tumour islands means that CEA could have little influence on interactions at the tumour/stroma interface.

Increasingly tumours other than adenocarcinomas are being recognised as expressing CEA related antigens. The development of monoclonal anti-CEA antibodies reacting with CEA on the surface of tumour cells has led to the successful immunolocalisation of CEA-expressing tumours and their metastases. Early work has suggested a role for CEA antibodies in tumour immunotherapy. This highlights the importance of localising and typing CEA related antigens in normal and neoplastic squamous tissues.

We wish to thank Mr Barrie Sims for the photomicrograph of the immunoblot. This work was funded by the United Birmingham Hospitals Endowment Fund (F08689).

References: