ORIGINAL ARTICLE

Characterisation of adherens and tight junctional molecules in normal animal larynx; determining a suitable model for studying molecular abnormalities in human laryngopharyngeal reflux

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Background: The disruption of intercellular junctions in the larynx is a pathological feature of laryngopharyngeal reflux (LPR). Good experimental models are necessary to gain greater insight into the molecular mechanisms and alterations that result from abnormal exposure of the laryngeal epithelium to acid refluxate.

Aims: To characterise laryngeal tissues from different species to determine the most suitable for use in experimental studies of LPR.

Methods: Human and non-human laryngeal tissues (mouse, rat, guinea pig, porcine, and rabbit) were studied. Histological characterisation was performed by light microscopy. The expression and subcellular localisation of adherens junctional molecules (E-cadherin and β catenin) was evaluated by immunohistochemistry, and tight junction molecules (occludin and zonula occludens 1 (ZO-1)) by western blotting. The ultrastructural features of porcine and human tissue were assessed by electron microscopy.

Results: Porcine tissue revealed both respiratory-type and stratified squamous epithelium, as seen in the human larynx. The expression and subcellular localisation of the E-cadherin–catenin complex was detected in all species except mouse and rat. The pattern of ZO-1 and occludin expression was preserved in all species.

Conclusion: The expression of intercellular junctional complexes in porcine epithelium is similar to that seen in humans. These results confirm the suitability of these species to study molecular mechanisms of LPR in an experimental system.

Laryngopharyngeal reflux (LPR) has gained increased recognition over recent years.^{1–8} Acid refluxate is known to contribute to the development of many otolaryngological symptoms and conditions, including laryngitis, sore throat, contact ulcers, and globus pharyngeus.⁹ However, to date, very little has been published in relation to the molecular mechanisms or alterations that result from abnormal exposure of acid refluxate on the laryngeal epithelium.

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The pathophysiology and molecular mechanisms of LPR are not completely understood. Both LPR and gastro-oesophageal reflux disease (GORD) are caused by acid and pepsin exposure.^{4–10} However, a direct association between mucosal findings and uncontrolled reflux in the larynx has not been established.¹¹ It is probable that acidified pepsin causes a break in the barrier function through an alteration in the intercellular junctional complex, as has been documented in the oesophagus.^{12–16} Indeed, we have recently shown that laryngeal E-cadherin expression is reduced or partially lost in laryngeal biopsies from patients with documented LPR.¹⁷

Intercellular junctions are composed of adherens junctions (E-cadherin–catenin complexes) and tight junction mole-

cules.^{18–23} Adherens junctions represent a large family of transmembrane glycoproteins (cadherins) responsible for calcium dependant intercellular adhesion and interaction.^{24–27} In particular, they are important for processes such as migration and restitution after damage. The tight junction complex consists of several cytoplasmic and membrane proteins, including zonula occludens 1 (ZO-1), and occludin, and is localised to the boundary of the apical and basolateral plasma membrane domains of epithelial cells.^{28–34} The tight junction complex forms a primary barrier to the passage of solutes through the paracellular space.

Good animal models are essential aids in the study of laryngeal pathophysiology. Concern exists, however, regarding the suitability of models because of both gross anatomical and histological differences to the human larynx. The aim of our study was to characterise the intercellular adhesion molecules in the laryngeal epithelium of several species and compare them with those seen in humans to identify the most suitable animal model for experimental studies on laryngeal reflux.

MATERIALS AND METHODS

Tissue specimens

Laryngeal tissue from non-human species (mouse, rat, guinea pig, pig, and rabbit) was obtained from the department of veterinary pathology, Langford, UK. Anonymised human laryngeal tissue samples were obtained from the

Abbreviations: GORD, gastro-oesophageal reflux disease; LPR, laryngopharyngeal reflux; ZO-1, zonula occludens 1

Table 1 Antibodies and their working dilutions

Antigen specificity	Antibody type	Source	Working dilution
E-cadherin	Mouse monoclonal	Transduction Laboratories	1/1000
β Catenin	Mouse monoclonal	Transduction Laboratories	1/1000
Occludin	Mouse monoclonal	Zymed	1/1000
ZO-1	Rabbit polyclonal	Zymed	1/1000
ZO-1, zonula occludens 1.			

department of otolaryngology at Southmead Hospital, Bristol. Approval was obtained from the local research ethics committee (reference 023/03).

Preparation of microscopic specimens

Specimens for light microscopy were immersion fixed in 10% neutral buffered formalin, dehydrated with ethanol, cleared with xylene, and embedded in paraffin wax. Transverse sections (4 μ m) were prepared and affixed to poly-L-lysine coated slides (Sigma, Poole, Dorset, UK). Haematoxylin and eosin staining was used to localise tissue structures.

For electron microscopy, laryngeal sections were selected under a dissecting microscope; these sections were cut into rectangular pieces and immersed in 2% glutaraldehyde in 0.1M sodium cacodylate buffer (pH 7.2) for two hours. The tissue was postfixed in 1% OsO_4 in 0.1M sodium cacodylate buffer (pH 7.2) for one hour, rinsed in 0.1M sodium cacodylate three times (10 minutes each), and deionised water (10 minutes). Specimens were then dehydrated in an ascending series of ethanol, replaced with propylene oxide, and flat embedded in epoxy resin for 72 hours at 60°C. Specific regions of the laryngeal tissue were located and identified using semithin sections (about 1 μ m) stained with 0.5% toluidine blue in 0.5% borate. Thin sections (0.1 μ m) were cut, stained with 3% uranyl acetate and lead citrate, and observed with a Phillips C200 electron microscope.

Monoclonal antibodies

Mouse monoclonal antibodies specific for E-cadherin and β catenin were purchased from BD Transduction Laboratories (distributed by Affinity, Exeter, UK) and antibodies to occludin from Zymed (Cambridge Biosciences, Cambridge, UK). Table 1 provides details of the antibodies used.

Immunohistochemistry

For immunohistochemistry, serial sections (4 μ m) were dewaxed, rehydrated, and incubated with the previously validated mouse monoclonal antibodies to E-cadherin, β catenin, occludin, and ZO-1 for one hour (table 1). Antigen retrieval was carried out for each antibody (micro-waving in 0.01M citrate buffer). Endogenous peroxidase activity was blocked with 3% hydrogen peroxide for 15 minutes. Antibody-antigen binding was detected using biotinylated goat and rabbit antimouse immunoglobulin (Dako, Glostrup, Denmark), followed by peroxidase conjugated streptavidin-biotin complex (Dako). Sites of peroxidase activity were demonstrated using 3,3'-diaminobenzidine (DAB; Sigma) for up to 10 minutes. The primary antibody was omitted in the negative control sections. Sections were examined for nuclear, cytoplasmic, and membranous expression by two independent observers (GG, MP). Expression and subcellular localisation (membranous, cytoplasmic, or nuclear) was evaluated in stratified squamous epithelium and respiratory-type epithelium).

Figure 1 Morphological characterisation of human, rat, and mouse laryngeal mucosal tissue by haematoxylin and eosin staining. (A) In the human larynx, the surface epithelial cell layer consists of ciliated columnar epithelial cells. Vascular channels and scattered chronic inflammatory cells are seen in the subepithelial fibroconnective tissue. The squamous epithelium of the laryngeal mucosa is largely non-keratinising and differs from rat sections (B). Respiratory-type epithelium was not seen in (B) rat or (C) mouse laryngeal tissue.

Protein preparation from tissue samples and western blot procedure

Tissue preparation

Protein extract was obtained by homogenisation of the tissue in lysis buffer containing 7M urea, 0.1M dithiothreitol, 0.05% Triton $\times$ 100, 25mM NaCl, and 20mM Hepes, pH 7.6. The extract was left on ice for 30 minutes and centrifuged at 10 000 $\times g$ for 15 minutes at 4°C. The pellet was discarded and the supernatant snap frozen and stored at -80°C .

Proteins were resolved on 7.5% polyacrylamide gels and transferred to Immobilon P polyvinylidene difluoride membranes (Millipore, Watford, Hertfordshire, UK). Each lane of this gel received 80 μ g of protein. After electrophoresis and transfer to membrane (Bio-Rad, Hercules, California, USA) by electroblotting, blots were probed with specific primary and secondary antibodies. All antibodies were diluted 1/1000. Blots were subsequently probed with anti- α tubulin (Sigma, Poole, Dorset, UK) to demonstrate equal sample loading. Detection was achieved using a horseradish peroxidase

Figure 2 E-cadherin immunostaining in laryngeal epithelium from (A) human, (B) mouse, and (C) rat.

conjugated antimouse secondary antibody. An enhanced chemiluminescence detection system (Amersham, Arlington Heights, Illinois, USA) was used for signal development, according to the manufacturer's protocol.

RESULTS

Histological characterisation of the laryngeal epithelium (haematoxylin and eosin sections)

Laryngeal epithelium from pig, guinea pig, and rabbit primarily consisted of three predominant cell types: ciliated, secretory (mucin producing), and stratified squamous epithelial cells similar to those found in the human larynx. Rat and mouse larynx showed alterations in tissue structure compared with the human tissue; namely, the squamous epithelium was largely keratinised in rat sections and respiratory-type epithelium was not identified in either rat or mouse tissue (fig 1).

Adherens and tight junction molecule expression and subcellular localisation

We analysed the expression and subcellular localisation of E-cadherin and β catenin in human and non-human larynx. In human larynx, E-cadherin was expressed over the entire intercellular membrane of the epithelial cells (fig 2A). The

Figure 3 Immunostaining for β catenin in laryngeal epithelium from (A) human, (B) mouse, and (C) rat.

Figure 4 Western blot analysis of E-cadherin and β catenin. Normal laryngeal tissue from (A) human, (B) pig, (C) guinea pig, (D) rat, (E) mouse, and (F) rabbit; α tubulin was used as a loading control. Note the presence of low intensity bands for mouse and rat tissue compared with the other species.

distribution of this molecule in the stratified squamous epithelium decreased with maturation and expression was undetectable in the superficial layers (fig 2A). The staining pattern of E-cadherin was also membranous in all the non-human laryngeal samples studied. However, compared with that seen in humans, the mouse and rat showed weaker immunostaining for this complex (fig 2B, C). In all sections studied, β catenin was present mainly in the membranous and to a lesser extent in the cytoplasmic cellular compartments. As

Figure 5 Western blotting analysis of zonula occludens 1 (ZO-1) and occludin. Normal laryngeal tissue from (A) human, (B) pig, (C) guinea pig, (D) rat, (E) mouse, and (F) rabbit; α tubulin was used as a loading control.

was seen for E-cadherin, immunoreactivity for β catenin was less intense in rat and mouse laryngeal samples (fig 3). To confirm these results, we evaluated the total intracellular protein content of E-cadherin- β catenin by western blotting. Indeed, we found low protein expression of both molecules in the samples from mouse and rat (fig 4).

Tight junction expression

We analysed the expression of the two tight junction molecules occludin and ZO-1 in human, mouse, rat, guinea pig, rabbit, and porcine tissue samples (fig 5). These molecules were expressed in all non-human species studied. In contrast to the E-cadherin-catenin complex, there was no difference in the total protein concentrations of occludin and ZO-1 between non-human and human tissues, as shown by western blotting.

Ultrastructural study of human and porcine laryngeal epithelium

Results obtained from histological and protein cellular expression studies clearly indicate that the porcine larynx is the most similar to the human larynx. For this reason, we investigated the ultrastructure of the porcine larynx by electron microscopy (fig 6A,B). We found stratified squamous and pseudostratified ciliated epithelium in the porcine larynx, similar to that found in human laryngeal sections. We noted a similar pattern of change in epithelial cell type from stratified squamous epithelial cells (supraglottic region) to pseudostratified, ciliated cells (subglottic region), as has been seen in human laryngeal tissue (fig 6C,D). The zonula adherens, zonula occludens, and desmosomal regions were similar in porcine and human tissue. Table 2 summarises the ultrastructural findings.

DISCUSSION

Our study investigated the anatomical and histopathological differences between the larynx in animals (pig, rabbit, guinea pig, rat, and mouse) and humans, with the aim of using the results to determine a suitable experimental model to study LPR. In this comparative study, we compared laryngeal anatomy and function of human and non-human species. We paid particular attention to essential adhesion structures (cadherin/catenins, ZO-1, and occludin) that may act as targets of acid refluxate.¹⁷ Acid and pepsin have been shown to break down the barrier function and epithelial permeability through perturbation of the junctional complex.³⁵ Both the protection and functional activity of the laryngeal epithelium rely upon structurally distinct cell categories,

Figure 6 (A) Transmission electron micrograph of the apical region of ciliated columnar and goblet cells from the vocal fold of human larynx tissue (original magnification, $\times 5200$). (B) Cell membranes of adjacent ciliated cells closely apposed at the junctional complex (arrow) (original magnification, $\times 11\,600$). (C) Transmission electron micrograph of the apical region of columnar epithelial and goblet cells from the porcine larynx (original magnification, $\times 5200$). (D) Cell membranes of adjacent cells closely apposed at the tight junctional complex (original magnification, $\times 11\,600$).

Table 2 Summary of ultrastructural results

Species	Laryngeal histology	E-cadherin	β Catenin	Occludin	ZO-1
Human	Stratified squamous epithelium; respiratory-type epithelium	++ (W)	++ (W)	(W)	(W)
Pig	Stratified squamous epithelium; respiratory-type epithelium	++ (W)	++ (W)	(W)	(W)
Guinea pig	Stratified squamous epithelium	+	+	(W)	(W)
Rat	Stratified squamous epithelium	+	+	(W)	(W)
Mouse	Stratified squamous epithelium	+	+	(W)	(W)
Rabbit	Stratified squamous epithelia; respiratory-type epithelium	++ (W)	++ (W)	(W)	(W)

E-cadherin and β catenin immunohistochemistry was scored using a semiquantitative scoring system: +, weak expression; ++, moderate expression as assessed by immunohistochemistry. (W), presence of protein as confirmed by western blot analysis.
ZO-1, zonula occludens 1.

and adherens junction and tight junction molecules play important roles.

E-cadherin plays an essential role in the structural and functional organisation of cells in tissues and organs of multicellular organisms. Sections from all non-human species showed uniform membranous E-cadherin staining patterns, similar to that seen in the normal human larynx. However we found differences in the total protein concentrations of E-cadherin, which was present in very low amounts in mouse and rat tissue. E-cadherin is able to bind the cytoskeleton through catenins. The presence of these catenins is therefore necessary for the full adhesive function of the cell. In the specimens examined, β catenin was localised uniformly at the intercellular junctions of the epithelial cells at the apical-lateral cell membrane compartment. Similar to E-cadherin, we found reduced protein expression of β catenin in rat and mouse tissue, as demonstrated by western blotting. It is possible that our findings may relate to specific antibody specificity to the different species investigated.

“Our findings indicate that from an ultrastructural, anatomical, and morphological perspective, the porcine larynx is a superior model for research into epithelial damage occurring during laryngopharyngeal reflux”

To our knowledge, there are no published data regarding the expression of tight junction molecules in the larynx. Occludin and ZO-1 are two integral membrane components that form tight junctions, especially in monolayers of epithelium derived tissues. In our study, we showed that in all species considered, ZO-1 and occludin were normally expressed and localised at the cell-cell contacts in the apical compartment. We confirmed the presence of these molecules by electron microscopy.

Our findings indicate that from an ultrastructural, anatomical, and morphological perspective, the porcine larynx is a superior model for research into epithelial damage occurring during LPR. We found that porcine tissue was the most comparable to human tissue because of the pattern of E-cadherin/catenin expression and the subcellular localisation.

Take home messages

- The expression of intercellular junctional complexes in porcine laryngeal epithelium is similar to that seen in humans
- The large size of the porcine larynx also offers a unique advantage over other models
- Thus, the porcine larynx is a suitable model for research into epithelial damage occurring during laryngopharyngeal reflux

The porcine larynx appears to be the most suitable to use in the development of an experimental model, in terms of practical use and similarity in tissue composition. The large size of the porcine larynx also offers a unique advantage over other models. Consequently, this animal model allows for the study of the common sites of reflux action seen in patients with LPR (supraglottic region and subglottic regions) within the same tissue sample.

Many different factors must be considered before selecting a suitable experimental model, in which the results can be translated to the clinical setting. Studying the pathological changes in the intercellular adhesion complexes after acid pepsin exposure could better characterise the molecular mechanisms involved in LPR. This aspect is currently under investigation in our laboratories, but additional studies are needed to determine the specific role of these ubiquitous molecules as clinical markers in the diagnosis of laryngeal reflux.

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