Telomerase activity detected in oral lichen planus by RNA in situ hybridisation: not a marker for malignant transformation

C O’Flatharta, M Leader, E Kay, S R Flint, M Toner, W Robertson, M J E M F Mabruk

Background: Oral lichen planus (OLP) is a chronic inflammatory condition. Clinically, it is characterised by the presence of a white lace-like lesion on the buccal mucosa, tongue, and gingivae, with erosions and ulceration. The World Health Organisation considers OLP to be a premalignant condition.

Aims: To investigate expression of the telomerase RNA component (hTR) in OLP compared with normal control buccal mucosa and to assess the possibility of using hTR expression as a marker for malignant transformation in OLP.

Methods: hTR expression was analysed in 40 cases of OLP and 18 normal control buccal mucosa samples using an RNA in situ hybridisation approach.

Results: Strong hTR RNA expression was seen in the basal, suprabasal, and to a lesser extent in the upper epithelial layers in 36 of the 40 OLP lesions examined. Infiltrating subepithelial lymphocytes in OLP were also shown to express hTR RNA. Weak hTR RNA expression was seen in seven of the 18 normal control buccal mucosa specimens, with expression confined exclusively to the basal layer of the epithelium and absent in the suprabasal and upper layers.

Conclusion: The telomerase RNA component hTR is found to be highly expressed in the epithelium of non-dysplastic OLP lesions. It is possible that this high expression is related to the increased cellular proliferation seen in OLP lesions rather than being an indicator of susceptibility to malignancy. Thus, hTR RNA expression may not be a suitable marker for predicting malignant transformation in OLP.

Telomerase is a specialised ribonucleoprotein polymerase composed of RNA and protein components and is thought to compensate for the loss of telomere during cell division. The hTR RNA moiety of telomerase has been cloned recently and its presence was shown to be crucial for telomerase activity. This integral RNA template element (hTR) directs the synthesis of telomeric repeats at chromosomal ends. Among the protein components are the human telomerase reverse transcriptase (hTERT, hEST2), which is thought to be the catalytic subunit of the enzyme, and human telomerase protein 1.

As one would expect, telomerase activity has been detected in immortalised and cancer cell lines, germ line cells, and most human malignancies, including head and neck squamous cell carcinomas.

Recent findings suggest that the activation of telomerase is a crucial step in the continuous proliferation of cancer cells, which allows them to attain immortality and therefore may be considered as a marker for tumour progression.

Telomerase activity is thought to be minimal or undetectable in normal differentiated somatic cells. However, recent

Abbreviations: hTR, RNA template component of human telomerase; OLP, oral lichen planus; PCR, polymerase chain reaction
published reports show that telomerase is expressed in a variety of regenerative tissues and may be associated with cellular proliferation. Low amounts of telomerase activity have been detected in studies including normal oral mucosa, normal human epithelial cells and cell cultures, blood, endometrial cells, and human hair follicles. A recent report demonstrated telomerase activity in OLP using the in vitro telomere repeat amplification protocol (TRAP) assay. However, the TRAP assay has several drawbacks. First, it does not allow for in situ analysis of telomerase activity. Second, the TRAP assay uses fresh or frozen tissue extracts and cannot identify which cells in tissue samples exhibit telomerase activity so that the precise contribution of infiltrating lymphocytes cannot be determined using this technique.

Therefore, in our present study, and for the first time, we have investigated the use of an RNA in situ hybridisation approach to elucidate the patterns of hTR RNA expression in OLP and normal buccal mucosa biopsies obtained from patients attending the Dublin Dental Hospital, Ireland.

### MATERIALS AND METHODS

#### Patients

Fifty eight formalin fixed, paraffin wax embedded tissue specimens were obtained from the Departments of Pathology, Beaumont Hospital and St James’s Hospital, Dublin, Ireland. These specimens consisted of 40 histologically confirmed non-dysplastic OLP tissue biopsies and 18 normal control buccal mucosa tissue biopsies without clinical evidence of OLP. Medical records and haematoxylin and eosin slides of the biopsies were reviewed by an oral pathologist (MT) to confirm the histological diagnosis.

The OLP group consisted of 11 men and 29 women, with a mean age of 48, ranging from 19 to 75 years (table 1). Twenty seven of the OLP specimens were taken from the buccal mucosa, four from the tongue, and one from each of the following: labial mucosa, maxillary alveolus, alveolar ridge, lower lip, and gingiva. Information on the biopsy site was unavailable for four patients (table 1). None of the patients in the study group went on to develop oral cancer during this time.

A lichenoid reactions, topographically related to buccal amalgam fillings, were excluded. Positive control normal testis tissue was analysed in each in situ hybridisation experiment.

#### In situ hybridisation

The RNA in situ hybridisation technique was applied to OLP and normal control buccal mucosal biopsies. All chemicals were supplied by Sigma-Aldrich (Poole, Dorset, UK) unless otherwise stated.

### Table 1: Relation between clinical appearance, biopsy site, sex, age, and expression of the RNA template component of telomerase in oral lichen planus tissue samples

<table>
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<tr>
<th>No.</th>
<th>Telomerase expression</th>
<th>Clinical appearance</th>
<th>Age</th>
<th>Sex</th>
<th>Biopsy site</th>
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<tr>
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<td>Labial mucosa</td>
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<td>BM</td>
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<tr>
<td>3</td>
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<td>RP</td>
<td>45</td>
<td>F</td>
<td>BM</td>
</tr>
<tr>
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<td>61</td>
<td>M</td>
<td>N/A</td>
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<td>60</td>
<td>F</td>
<td>BM</td>
</tr>
<tr>
<td>6</td>
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<td>RP</td>
<td>66</td>
<td>F</td>
<td>Tongue</td>
</tr>
<tr>
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<td>BM</td>
</tr>
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<td>BM</td>
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<tr>
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</tr>
<tr>
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<td>BM</td>
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<td>F</td>
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<td>BM</td>
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<td>F</td>
<td>BM</td>
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<tr>
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<td>F</td>
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<td>F</td>
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<td>M</td>
<td>BM</td>
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<tr>
<td>39</td>
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<td>Erosive reticular</td>
<td>N/A</td>
<td>M</td>
<td>N/A</td>
</tr>
<tr>
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<td>Positive</td>
<td>Erosive</td>
<td>N/A</td>
<td>M</td>
<td>N/A</td>
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</tbody>
</table>

Alv ridge, alveolar ridge; BM, buccal mucosa; N/A, data not available; RP, reticular papular.
the telomerase gene (TRC3R/TRC3F), subcloned into a PCR script vector containing both a T3 and a T7 promoter.

PCR amplification

PCR was performed with self designed primers: primer AB2, 5'-GTT TTC CCA GTC ACG ACG-3' (sense) and primer AB1, 5'-GAA TTG TGA GCG GAT AAC-3' (antisense). These primers were designed to amplify a region containing the insert of interest (in this case the TRC3R/TRC3F fragment of human telomerase flanked by the T3 and T7 promoters).

The PCR was conducted in a 50 µl reaction volume. Each reaction contained 1× PCR buffer (10mM Tris/HCl (pH 9.0 at 25°C), 50mM KCl, and 0.1% Triton X-100), 1mM MgCl₂, 0.2mM of each dNTP, and 5 U of Taq DNA polymerase (Promega, Wisconsin, USA). Initial denaturation at 95°C for five minutes was followed by 30 cycles of denaturation at 94°C for 45 seconds, annealing at 37°C for 45 seconds, and extension at 72°C for one minute in a programmable DNA thermal cycler (Hybaid, Teddington, UK). A five minute extension step was carried out after the last cycle to ensure complete polymerisation. For the electrophoretic analysis of amplified DNA, a 10 µl aliquot of PCR product was electrophoresed on a 1% agarose gel and visualised by ethidium bromide staining.

Probe labelling

Probe labelling was performed as described by the manufacturer (Roche Molecular Biochemicals, Lewes, East Sussex, UK), using T3 RNA polymerase and T7 RNA polymerase to create the antisense and sense probes, respectively.

Figure 1  Haematoxylin and eosin staining showing the characteristic histological features of oral lichen planus.

Figure 2  In situ detection of telomerase RNA gene expression in oral lichen planus biopsies. Expression of the telomerase RNA gene (hTR) was detected by in situ hybridisation with a riboprobe labelled with digoxigenin. Using an antisense probe, telomerase RNA expression appeared as blue/purple cytoplasmic staining in [A] the basal and suprabasal layers of epithelium, and [B] the subepithelial infiltrating lymphocytes (arrowheads). (C) No staining was observed using the sense RNA probe.
Probe precipitation
The labelled RNA was precipitated by the addition of 4M LiCl, 100% ethanol (−20°C) and glycogen, and the mix was then stored overnight at −20°C. Each reaction mix was subsequently centrifuged at 12,000 × g for 15 minutes. The resulting pellet was washed with 70% ethanol (−20°C) and then air dried. The dried pellets were resuspended in sterile RNase free water. The probe was further purified by passing it through a Nick Column (Amersham Pharmacia Biotech AB, Uppsala, Sweden), as described by the manufacturer. The integrity of the RNA probe was verified by electrophoresis on a 1% BSSE gel (50 mM boric acid, 5mM sodium tetraborate, 10mM sodium sulphate, 0.1mM EDTA) and visualised by ethidium bromide staining.

Digoxigenin labelled RNA in situ hybridisation
Sections of formalin fixed, paraffin wax embedded normal testes were used as a positive control for the detection of telomerase expression. The sections of testes were cut at a thickness of 4 μm on a microtome (Microm HM325; Walldorf, Germany) and baked on to silanised slides at 57°C overnight.

In situ hybridisation
The prehybridisation and post hybridisation treatments used were essentially the same as described previously by Soder et al., except that we used a digoxigenin labelling approach.

Visualisation of digoxigenin labelled probes
This was carried out as described by the manufacturer using the digoxigenin nucleic acid detection kit (Boehringer Mannheim/Roche Molecular Biochemicals).

Statistical analysis
Statistical analysis was carried out using Fisher’s exact test.

RESULTS
Histopathology
The biopsies showed features of OLP with varying degrees of keratosis. In some, the epithelium was hypertrophic, whereas others were atrophic. All had a lymphocytic infiltrate with exocytosis and basal layer liquefactive necrosis, some containing numerous Civatte bodies. Some of the biopsies included areas of ulceration and most of these contained plasma cells focally within the infiltrate, adjacent to areas of classic OLP. In a few cases, there was a subtle subepithelial split but no obvious bulla formation. None of the cases showed dysplasia. Figure 1 shows a typical example of OLP.

Detection of hTR RNA expression by in situ hybridisation
The telomerase antisense probe (T7) gave clear blue/purple cytoplasmic staining in the epithelial cells of the seminiferous tubules of the testis (positive control specimen). Using the telomerase sense probe (T3) no staining of any kind was seen in the epithelial cells of the seminiferous tubules of the testis with each in situ hybridisation experiment performed, thereby controlling for the specificity of each experiment.

The in situ hybridisation expression of hTR was detected in the epithelium of 36 of the 40 OLP lesions examined using the antisense RNA probe.

The expression patterns of hTR RNA were heterogeneous and confined to the epithelial cytoplasm, with strong expression of hTR RNA in the basal and suprabasal layers of all hTR RNA positive OLP cases (fig 2A).

In five of the 36 positive cases the more superficial prickle cells were positive for hTR RNA.

Expression of hTR RNA was also detected in a high proportion of the sublesional infiltrating lymphocytes in 35 of the 40 OLP cases (fig 2B).

DISCUSSION
In most human tumours and tumour cell lines, the telomere maintenance mechanism involves the activity of telomerase. RNA in situ hybridisation experiments have previously shown the telomerase RNA gene (hTR) to be expressed in 81.6–100% of oral squamous cell carcinomas and also in oral hairy leukoplakia. Using the TRAP assay, telomerase activity has been detected in 75–100% of oral squamous cell carcinomas and 43.4–86% of oral hairy leukoplakias.

A recent report published using the TRAP assay (an in vitro PCR based method) detected telomerase activity in 70% of OLP tissues investigated. This TRAP assay has several disadvantages. First, the TRAP assay does not allow the visualisation of the exact cell types in a tissue sample that are expressing telomerase activity. Second, fresh or frozen tissue is required for the technique. Third, the TRAP assay is complicated and can be affected by many factors. For example, in the case of oral lesions, false positive results may be the result of positivity in infiltrating lymphocytes and contaminating factors such as bacteria and blood. In contrast, the in situ hybridisation approach is more informative because it has the ability to detect and localise the expression of hTR in formalin fixed, paraffin wax embedded archival tissue.

“We have applied a novel approach for generating an RNA probe for the in situ hybridisation technique”

In our present study, we have applied a novel approach for generating an RNA probe for the in situ hybridisation
technique. This approach was developed in our laboratory to detect the expression of hTR in archival formalin fixed, paraffin wax embedded tissue.

We have applied this in situ approach to OLP biopsies for the first time. Telomerase hTR RNA expression was detected in the basal cells, suprabasal cells, and more superficially located prickle cells in 36 of the 40 OLP specimens examined.

Four of our OLP samples did not express hTR RNA. The study by Thongprasom et al, which used the TRAP assay, found that 30% of OLP samples did not exhibit telomerase activity.

In addition, we detected low amounts of hTR expression in the basal layer of seven of the 18 normal control buccal mucosa tissue samples. Expression of hTR was considerably weaker than that seen in the OLP specimens and no expression was seen in the suprabasal or upper layers of the mucosa. Recent in situ hybridisation studies have shown hTR expression in the basal layers of normal oral epithelia, normal cervical epithelium, and epidermal basal cells from sun exposed skin.

Although telomerase activity has been shown to be absent in most normal somatic tissues, there have been reports of telomerase activity in cells with a high proliferation potential, such as normal oral squamous epithelium, haematopoietic stem cells and lymphocytes, cultured normal human epidermal keratinocytes, and mitotically active segments of human hair follicles. The detection of low amounts of telomerase activity in some histologically normal tissues probably reflects the presence of proliferating stem cells in these samples.

OLP is an immune mediated disorder characterised by the presence of a dense band-like inflammatory infiltrate at the epithelial/subepithelial interface. These activated lymphocytes may have telomerase activity and hTR expression. We examined each OLP specimen for the presence of hTR expression in the infiltrating lymphocytes. This in situ hybridisation technique detected hTR expression in the lymphocytes within the inflammatory infiltrate in OLP. hTR expression in the lymphocytic infiltrate corresponded with the presence of lymphocytes seen in the haematoxylin and eosin stained section for each OLP sample. Although lymphocytic hTR expression has not been detected in in situ hybridisation studies involving the oral mucosa, in situ studies on normal and cancerous cervical tissue and other non-neoplastic livers have revealed hTR expression localised to infiltrating lymphocytes.

In addition, moderate hTR expression was limited to the secondary lymphoid follicles (for example, lymph node and spleen) in embryonic tissues, whereas hTR expression was also shown to be present in the germinal centres of benign lymph nodes.

It has been shown previously that the treatment of immortalised human cell lines with antisense oligonucleotides for hTR limits their proliferative capacity. Recent studies have reported an association between hTR expression and the degree of malignancy in oral mucosal lesions, and a correlation between hTR expression and telomerase activity has been described in lung and cervical cancer.

Although telomerase was thought to be associated mainly with the malignant phenotype, evidence has accumulated that telomerase may be related to cellular proliferation. Telomerase activity has been reported in endometrial tissue and was shown to correlate with proliferation during the menstrual cycle. It has been shown previously that OLP has an increased proliferative capacity compared with other oral keratoses and normal mucosa. It is possible that the increased expression of the telomerase RNA component in OLP may be evidence of the hyperproliferative state, as shown by the determination of growth fraction.

“We hypothesise that hTR expression is associated with the high proliferation rate of OLP and does not necessar-

Take home messages

- The in situ hybridisation approach is more informative than the telomere repeat amplification protocol because it has the ability to detect and localise the expression of the telomerase RNA component (hTR) in formalin fixed, paraffin wax embedded archival tissue
- hTR is highly expressed in the epithelium of non-dysplastic oral lichen planus (OLP) lesions
- However, we think that this high expression is related to the increased cellular proliferation seen in OLP lesions rather than being an indicator of susceptibility to malignancy
- Therefore, hTR RNA may not be a suitable marker for predicting malignant transformation in OLP

The potential for malignant transformation in OLP is relatively low. Our study has shown that the telomerase RNA component is expressed in the suprabasal epithelium of 90% of the OLP samples examined. Other investigators have detected telomerase activity by TRAP assay in 70% of a smaller number of OLP tissues, although this last study did not include normal control tissue.

This and previous studies have demonstrated a relatively high expression of hTR and telomerase activity in OLP. We suggest that hTR expression is associated with the high proliferation rate of OLP rather than being an indicator of malignant transformation.

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