Original Article

DNA Fragmentation and mRNA Expression of Bcl-2, Bcl-xL, p53, p21, and HSP70 Genes in Nondysplastic and Dysplastic Oral Lichen Planus

Abstract

Background: Oral lichen planus (OLP) is a chronic inflammatory disease. Apoptosis of the basal keratinocytes is a causative factor for OLP pathogenesis, but the detailed mechanism of apoptosis among nondysplastic and dysplastic OLP lesions is yet unraveled. Aims: This study aims to evaluate the involvement of cellular DNA fragmentation and alteration in the expression of Bcl-2, Bcl-2, and B-cell lymphoma extra-large (Bcl-xL), p53, p21 and Heat shock protein 70 (HSP70) in nondysplastic and dysplastic OLP lesions. Materials and Methods: Untreated, fifteen OLP patients each with nondysplastic and dysplastic lesions were enrolled for this study. Their DNA fragmentation was analyzed by the agarose gel electrophoresis method. The mRNA expression of Bcl-2, Bcl-xL, p53, p21, and HSP70 were measured using semi-quantitative reverse transcription-polymerase chain reaction. Results: Elevated DNA fragmentations were found in dysplastic lesions compared to nondysplastic type. Significantly higher expression of Bcl-2, Bcl-xL, p53, and p21 was found in both types of OLP lesion compared to the control. Expression of Bcl-2 and Bcl-xL were significantly elevated in nondysplastic lesions, whereas significantly overexpression of p53 and p21 were found in dysplastic lesions. Anti-stress protein HSP70 was overtly expressed in dysplastic lesions compared to other groups. Conclusion: Reduced expression of Bcl-2 and Bcl-xL, with elevated DNA fragmentation, may be associated with increased apoptosis in dysplastic lesions which aids in the resolution of the chronic inflammatory process. Higher expression of p53 and p21 in dysplastic lesions reflects its malignant potentiality. Overexpression of HSP70 in dysplastic lesions is a useful marker for higher cellular stress.

Keywords: Apoptosis, DNA fragmentation, dysplasia, malignant potentiality, oral lichen planus

Introduction

Lichen planus is a chronic inflammatory disease condition that affects the skin and the oral mucus membrane. etiology and detailed pathogenesis of oral lichen planus (OLP) has not yet understood.[1] Clinically, OLP appears as white striations (Wickham's striae), white papules, white plaque, erythema, erosion, or blisters.[2] It has several subtypes including reticular. erosive, atrophic, papular. plaque-like, and bullous and some variants may co-exist.[3] Apoptosis is frequently found in OLP, but the mechanism yet to be clarified.[4] It was found that apoptosis prevents aneuploidy and other genetic alterations which are associated with the development and progression of precancerous lesions.[5] Among the other important markers of apoptosis, fragmentation is considered

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hallmark.^[6] OLP has a higher risk of malignant transformation compared to the cutaneous lesion, the reported range being 0%–12.5%.^[7] The presence of oral epithelial dysplasia is an important predictor for malignant transformation in oral premalignant disorders like OLP.^[8]

Expression of pro-and anti-apoptotic proteins is important to understand the nature of oral lesion and the process in disease progression.^[9] Among the others, proapoptotic Bcl-2 associated X protein, Bcl-2 associated agonist of cell death, Bcl-2-related ovarian killer, annexin, and anti-apoptotic Bcl-2 and B-cell lymphoma extra-large (Bcl-xL) having an important regulatory role in oral lesions formation.[10] Cellular stress produced by DNA-damaging agents can activate apoptosis by the interplay of pro- and anti-apoptotic Bel family proteins.[11] There is a strong correlation between the expression of p53 and Bcl-2 proteins and neoplastic

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transformation.^[12] Heat shock protein 70 (HSP70) is a major stress-inducible protein involved in the protection of cells from many apoptotic stimuli.^[13] HSP70 might be a promising molecule for controlling apoptosis.^[14]

DNA fragmentation and role of anti-apoptotic Bcl-2, Bcl-xL, cell cycle regulator p21, anti-stress HSP70 gene expression in pathophysiological alteration of nondysplastic and dysplastic OLP lesions formation has not yet looked into. Hence, the present study aims to analyze the DNA fragmentation and alteration in mRNA expression of Bcl-2, Bcl-xL, p53, p21, and HSP70 in nondysplastic and dysplastic OLP lesions. This study may be useful to identify the causative factors of nondysplastic and dysplastic OLP lesion formation.

Materials and Methods

Patient selection

Freshly diagnosed and untreated 15 OLP patients (without any dysplasia) and another 15 OLP patient (with mild-to-moderate dysplasia) aged 35-55 years were enrolled for this study. A total of 15 age- and sex-matched healthy individuals were recruited as control subjects. The patients were clinically and histologically diagnosed according to modified WHO criteria.[15] Punch biopsy technique was used to collect the OLP lesion tissue samples from the Department of Oral Medicine and Radiology, PMS College of Dental Science and Research, Kerala, India. Normal healthy samples were the discarded tissues from surgical treatments of the impacted tooth, where the normal mucosal tissue margins were trimmed (excised) and discarded to facilitate primary closure, which were collected from the aforesaid department. The data and tissue sample collection were approved by the Institutional Ethical Committee and written consent for publication was obtained.

Exclusion criteria

OLP patients who had undergone any treatment for the same, those previously with habits of chewing or smoking tobacco and those with a history of alcohol consumption were eliminated from this study. OLP patients or normal individuals having asthma, hypertension, diabetes, cardiac disorder, bleeding or clotting disorders, psychiatric illness, hepatitis, AIDS, or malignancy were not included in this study.

DNA fragmentation were analyzed by agarose gel electrophoresis method

25 mg tissue was placed in a sterile microcentrifuge tube, 180 μl PureLinkTM genomic digestion buffer, and 20 μl proteinase K. The mixture was incubated at 55°C with occasional vortexing until lysis was complete (1–4 h). The lysate was centrifuged at maximum speed for 3 min at room temperature and the supernatant was transferred to a new sterile tube. 20 μl RNase A was added to the lysate, mixed well and incubated at room temperature

for 2 min. 200 μ l PureLinkTM genomic lysis/binding buffer was added and mixed well by vortexing to yield a homogeneous solution. 200 μ l 96%–100% ethanol was added to the lysate and mixed well to obtain a homogeneous solution.

Determination of mRNA expression by two-step semi-quantitative reverse transcription-polymerase chain reaction

The mRNA expression of Bcl-2, Bcl-xL, HSP70, p53, p21, and GAPDH in OLP lesion groups and control tissue were studied by using reverse transcription-polymerase chain reaction (RT-PCR) method. Total RNA was isolated from oral tissue using the TRIzol reagent (Invitrogen, USA) according to the manufacture's instruction. RT-PCR was performed using primer designed specifically for the genes to be amplified. cDNA was synthesized using ThermoScriptTM kit (Invitrogen, USA). The synthesized cDNA was amplified using Platinum Taq DNA polymerase kit (Invitrogen, USA). The stained gel was observed using the E-gel imager system (Invitrogen, Life Technology, USA). Agarose gel pictures were analyzed by Image J software Table 1.

Statistical analysis

The results were expressed as mean \pm standard deviation and the statistical analyses were performed using one-way ANOVA and Tukey's *post hoc* test by SPSS version 18.0 (SPSS Inc., Chicago, IL, USA). P < 0.05 has been considered as statistically significant.

Results

A cross-sectional study was performed on 15 OLP patients with oral dysplasia and another 15 OLP patients without any dysplasia. The clinical appearance, site of OLP lesions, and type of dysplasia in OLP patients have been tabulated [Table 2]. The qualitative analysis of DNA fragmentation was used to detect apoptosis in both types of OLP lesions and healthy control. The mRNA expression of

Table 1: Gene specific forward and reverse primer sequences of apoptosis associated genes

Gene	Primer sequence
Bcl-2	Forward 5'-CCTGTGGATGACTGAGTACC-3'
	Reverse 5'-GAGACAGCCAGGAGAAATCA-3'
Bcl-xL	Forward 5'-GAGTTCATTCACTACCTGTTC-3'
	Reverse 5'-CTGGTGGTTGACTTTCTCTC-3'
p53	Forward 5'-CCACCATGAGCGCTGCTCA-3'
	Reverse 5'-GCAGGGGAGGGAGAGATG-3'
p21	Forward 5'-GCAGATCCACAGCGATATCC-3'
	Reverse 5'-CAACTGCTCACTGTCCACGG-3'
HSP70	Forward 5'-CCATGGTGCTGACCAAGATGAAG-3'
	Reverse 5'-CACCAGCGTCAATGGAGAGAACC-3'
GAPDH	Forward 5'-GAAGGTGAAGGTCGGAGTC-3'
	Reverse 5'-GAAGATGGTGATGGGATTTC-3'

Table 2: Basic characteristics of patients with non-dysplastic and dysplastic changes											
Nondysplastic OLP lesions						Dysplastic OLP lesions					
Case number	Gender	Age	Site of lesion	Clinical appearance	Gender	Age	Site of lesion	Clinical appearance	dysplasia		
1	Female	36	Buccal mucosa	Reticular	Female	48	Buccal mucosa	Erosive	Moderate		
2	Male	46	Buccal mucosa	Atrophic	Female	53	Buccal mucosa	Erosive	Mild		
3	Female	39	Lateral border of tongue	Reticular	Female	55	Buccal mucosa	Plaque	Mild		
4	Female	41	Buccal mucosa	Reticular	Female	44	Lateral border of tongue	Erosive	Moderate		
5	Female	52	Buccal mucosa	Atrophic	Female	38	Floor of mouth	Erosive	Mild		
6	Male	37	Floor of mouth	Atrophic	Male	42	Lateral border of tongue	Plaque	Mild		
7	Male	36	Buccal mucosa	Erosive	Male	54	Buccal mucosa	Erosive	Moderate		
8	Male	35	Lateral border of tongue	Reticular	Male	36	Buccal mucosa	Plaque	Mild		
9	Female	42	Buccal mucosa	Reticular	Female	45	Buccal mucosa	Erosive	Moderate		
10	Female	50	Buccal mucosa	Atrophic	Female	49	Lateral border of tongue	Plaque	Mild		
11	Female	47	Floor of mouth	Reticular	Male	37	Lateral border of tongue	Erosive	Moderate		
12	Male	44	Buccal mucosa	Reticular	Male	51	Buccal mucosa	Plaque	Mild		
13	Female	43	Buccal mucosa	Popular	Female	38	Lateral border of tongue	Plaque	Mild		
14	Female	51	Lateral border of tongue	Reticular	Male	53	Buccal mucosa	Erosive	Moderate		
15	Male	41	Floor of mouth	Erosive	Male	47	Buccal mucosa	Erosive	Mild		

OLP: Oral lichen planus

Bcl-2, Bcl-xL, p53, p21, and HSP70 were compared among nondysplastic, dysplastic OLP lesions and control samples.

The presence of saw-toothed rete ridge, Max-Joseph space, and inflammatory infiltration was identified in normal, mild, and moderate dysplastic OLP lesions. Among other identifying characters, hyperchromatic nuclei were commonly observed in mild and moderate dysplastic lesions. In addition, increased nucleocytoplasmic ratio and enlarged nucleoli were found in mild dysplastic lesions, whereas, mitotic figure and basal cell hyperplasia were seen in moderate dysplastic lesions.

DNA fragmentation is a classic feature of apoptosis, cellular DNA is broken down into smaller fragments by the enzymes which are activated under apoptotic stimuli. This DNA fragmentation produces a characteristic pattern that can be visualized by gel electrophoresis, termed as DNA laddering. Higher fragmented DNA patterns were observed in mild and moderate dysplastic OLP lesions as compared with nondysplastic and healthy controls.

The mRNA expression of Bcl-2 (P < 0.001), Bcl-xL (P < 0.0001), p53 (P < 0.0001), and p21 (P < 0.0001) significantly differed from the control, nondysplastic and dysplastic OLP lesion group patients. Significantly higher expression of Bcl-2 and Bcl-xL were found in OLP groups compared to the control, whereas, expression of both anti-apoptotic genes was significantly (P < 0.05) elevated in nondysplastic lesions compared to the dysplastic type. Expression of transcription factor p53 and cell cycle regulator p21 was significantly elevated in OLP groups compared to control. Significantly higher expression of p53 (P < 0.0001) and p21 (P < 0.006) were found in dysplastic lesions compared to nondysplastic OLP. Importantly, significant overexpression

HSP70 (P < 0.0001) was seen in dysplastic OLP lesions when compared with nondysplastic and control tissue.

Discussion

OLP is a T-cell-mediated autoimmune disease of unknown etiology. Apoptosis of basal keratinocytes of the oral epithelium initiates the pathogenesis OLP. [16] The stimulators for apoptosis in OLP are still unknown, in which auto-cytotoxic CD8+ T cells attack the basal epithelial cells and activate vital molecular mechanisms namely cell cycle arrest for DNA repair and induce cell senescence or apoptosis to eliminate cells with severely damaged DNA. [17] DNA fragmentation is the main characteristic feature of apoptosis and thus is used as an important marker of apoptosis. [18] In the present study, DNA fragmentations in dysplastic lesions were higher compared to nondysplastic and control type which is an indicator for the high rate of apoptosis in dysplastic lesions.

The growth of oral keratinocytes is regulated by a delicate balance between anti-apoptotic Bcl-2 that controls cell survival and p53 that controls cell death.[19] Alteration in the expression of these proteins is a strong indicator of the malignant transformation potential of a certain lesion.[20] The overexpression of the Bcl-2 and Bcl-xL was found to protect the cells from undergoing apoptosis in response to a number of apoptotic stimuli.[21] The detailed mechanism by which Bcl-2 inhibits apoptosis is still uncertain, it was suggested that Bcl-2 and Bcl-xL may inhibit mitochondrial cytochrome c translocation and simultaneously prevent caspase activation by which they prevent the apoptotic process.[22] Anti-apoptotic Bcl-2 and Bcl-xL protein were the first known molecular targets of mitochondrial p53.[23] Interestingly, p53 can participate in apoptosis induction by direct action on mitochondria. The apoptotic signals cause

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localization of p53 into the mitochondria to precede the release of cytochrome c and activation of procaspase-3. [24]

In the case of OLP, 92.9% Bcl-2 positive staining was found in inflammatory infiltration.^[25] The overexpression of Bcl-2 was reported in oral dysplastic lesions and it

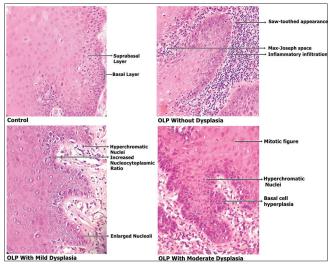


Figure 1: Histological images showing the presence of different characteristic features in nondysplastic and dysplastic lesions (H and E; ×40)

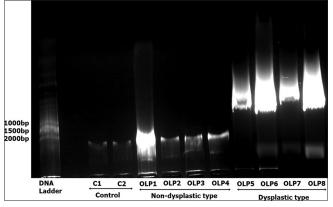


Figure 2: DNA fragmentation pattern in non-dysplastic and dysplastic OLP lesions compared with control

was suggested that Bcl-2 plays an important role in apoptosis and oral tumorigenesis. [26,27] Whereas the lack of expression of Bcl-2 in oral dysplasia was observed in other studies. [28,29] Importantly, dysregulation of the Bcl-2 gene expression in oral epithelial dysplasia may be an important causative factor for genetic aberrations in the progression of epithelial tumors. [9] In the present study higher expression of Bcl-2 and Bcl-xL are observed in all types of OLP lesions irrespective of OLP groups, indicating the apoptotic nature of OLP lesions. Comparatively, reduced expression of Bcl-2 and Bcl-xL in dysplastic lesions is an important indicator for a higher degree of apoptosis.

Different forms of oncogenic stress and DNA damage induce cell-cycle arrest, apoptosis, or senescence which may activate the tumor-suppressor p53 to protect cells from malignant transformation. This cellular accumulation of p53 protein can activate multiple other target genes which are required for tumor suppression. The cyclin-dependent kinase inhibitor/p21 is one of them, whose promoter contains two p53-binding sites, where at least one of them is essential for p53 binding after DNA damage. This binding inhibits apoptosis by causing cell cycle arrest and DNA repair. Later, the p21-arrested cells may undergo apoptosis by the activation of pro-apoptotic genes.

Previous studies have shown overexpression of p53 in OLP plays a regulatory role in apoptosis. [34] Elevated expression of p53 and p21 was observed in OLP patients when compared to normal but this did not differ from mild epithelial dysplasia may indicating the precancerous potentiality of OLP lesions. [20,35] In the present study, mRNA expression of p53 and p21 were significantly elevated in both types of OLP patients compared to healthy individuals. Higher expression of p53 and p21 in dysplastic lesion compared with nondysplastic may play a critical role in apoptosis signaling as well as malignant transformation.

Most HSPs are rapidly overexpressed in response to cellular injuries including genotoxic stress. HSPs protect cells from apoptosis by preventing mitochondrial apoptosome formation and caspase activation. [36] HSPs promote tumorigenesis by suppressing

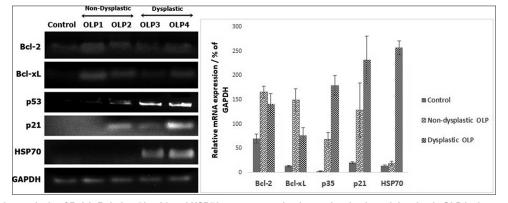


Figure 3: Comparative analysis of BcI-2, BcI-xL, p53, p21 and HSP70 gene expression in nondysplastic and dysplastic OLP lesion compared to control

apoptosis.^[37] Increased expression of HSP70 is a marker for the presence of epithelial dysplasia or epithelial malignant transformation. ^[38,39] Upregulation of HSP70 was noted in premalignant lesions and oral squamous cell carcinoma. ^[40] Altered expression of HSP70 has been reported in OLP and oral leucoplakia, with the highest intensity in severe oral dysplasia which suggested that HSP70 may be an objective marker for the identification of epithelial dysplasia. ^[41,42] Similar results were found in our study, where HSP70 was overtly expressed in dysplastic lesions compared to nondysplastic type. This may be a prime indicator for higher cellular stress and the chance of malignant transformation in dysplastic OLP lesions compared to the nondysplastic type.

Alteration of multiple signaling pathways may cause deregulated apoptosis which is an important contributor for malignant transformation. This is a mechanism of immune evasion which causes delayed apoptosis and prolongs inflammatory cell survival. The detailed pathogenic mechanism of OLP is partially understood, most importantly the initial events in lesion formation and mechanism of keratinocytes apoptosis in OLP need to be addressed. The comparative analysis of DNA fragmentation and expression of HSP70 in nondysplastic and dysplastic OLP lesions have not yet looked into, so this study may provide critical information about the pathogenesis mechanism.

Conclusion

Apoptosis is an important biological process involved in physiology and pathology. Nondysplastic OLP lesions have potency to transform a dysplastic lesion and the dysplastic lesions having higher malignant potentiality but the detailed mechanism yet to be understood. In the present study, we have tried to evaluate the causative factors for the dysplastic and nondysplastic OLP lesion formation, which having an important role pathogenesis mechanism. Intracellular stress signaling causes high DNA fragmentation which activates apoptotic pathways in OLP lesions. Elevated DNA fragmentation and reduced expression of Bcl-2 and Bcl-xL in the dysplastic lesion is an indicator of the high rate of apoptosis when compared with nondysplastic type. Overexpression of p53, p21, and HSP70 in dysplastic OLP lesions have regulatory role in apoptosis signaling as well as in malignant transformation.

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Conflicts of interest

There are no conflicts of interest.

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