INTRODUCTION

Oral submucous fibrosis (OSMF) is a chronic, insidious and progressive oral mucosal disease that primarily affects any part of the oral cavity. It is characterized by a juxta-epithelial inflammatory reaction followed by progressive fibrosis of the lamina propria and the underlying submucosal layer, with associated epithelial atrophy. Although the etiology of OSMF is obscure, evidence has shown that it is a precancerous disorder related to the habit of chewing areca nut, either alone or as a component of betel quid. OSMF carries a high risk of transition to oral cancer. In an epidemiologic study in India, the malignant transformation rate was 7.6% to 12 over a period of 17 years.[2]

Cancer induction is a multi-stage, multi-step process and includes multiple molecular and cellular events to transform a normal cell into a malignant neoplastic cell. However, three general steps can be identified in carcinogenesis; initiation, promotion and progression.[3]
Recently it has been proposed that oral cancers arising in OSF constitute a clinicopathologically distinct disease, the differences of which is believed to arise from differential mechanisms of areca nut carcinogenesis. A study recognized that most of these patients are younger males with better prognostic factors such as better grade of tumor differentiation, a lesser incidence of nodal metastases and extracapsular spread.[4] Another retrospective study has reported contradictory data. They state that oral squamous cell carcinoma (OSCC) can originate from OSF and is clinically more invasive and also exhibits higher metastasis and recurrence rate than OSCC that has not originated from OSF.[5]

We hypothesize that progression of OSMF and malignant transformation in the background of fibrosis mediates via hypoxia inducible factor-1α (HIF-1α) either by up- or down-regulation of various such molecules.

**MATERIALS AND METHODS**

Twenty blocks of formalin-fixed and paraffin-embedded tissues of each OSMF, OSCC and OSCC with OSMF cases were randomly selected from the archives of the Department of Oral Pathology and Microbiology. All 60 cases and 10 normal oral mucosal biopsies were used to investigate the expression levels of HIF-1α by immunohistochemistry. Ethical approval and informed consent from patients were obtained for this study.

**Grading and stage of the disease**

Routine hematoxylin and eosin stained sections were prepared from all the cases. The histological grading of OSMF was carried out. Also, the squamous cell carcinomas were graded as well, moderate and poor according to the Broder’s and Bryne’s grading system[6-8] [Figure 1].

**Immunohistochemistry**

Formalin-fixed paraffin-embedded tissue sections were cut to 5 μm thickness. Silane coated slides were used for the proper adherence of tissue sections to the glass slides. The immunohistochemical staining for HIF-1α using Universal Immuno-Enzyme Polymer Technique was carried out in the Department of Oral Pathology and Microbiology. Sections were hydrated with increased grades of alcohol and brought to distilled water and treated with hydrogen peroxide to eliminate endogenous peroxidase activity. Then antigen retrieval with tri-sodium citrate for HIF-1α was carried out. The tissue was incubated sequentially with:

- Primary antibodies, that is, HIF-1α
- DAKO Envision system HRP labeled polymer detection system
- 3,3, Diaminobenzidine substrate solution – This results in formation of a colored precipitate at the tissue antigen sites. Visualization was aided by counterstaining with hematoxylin.

**Assessment of the hypoxia inducible factor-1α staining intensity**

Section was considered either as negative or positive according to the absence or presence of brown staining in epithelial or stromal cells. The positive cases were graded into three categories depending on the intensity of staining. Homogenous dark brown staining was considered as strong (+++) and light faint staining was considered as mild (+) and cases in between the two extremes were graded into the moderate (+++) category[2] [Figure 2].

**Assessment of mean blood vessel density**

The most intense vascular area was found by light microscope in ×40. Five such fields were selected under ×40 for each slide, with slide moving in clockwise direction. The area representing vascular tissue in the three digital images were imported to image analysis software and the area counted. This indicated vascular tissue area. The area representing total tissue in the three digital images were imported to image analysis software and the area counted. This indicated total tissue area. Mean vessel density (MVD): The ratio of the vascular tissue area to total tissue area in the digital images imported to image analyzer software.
The number of blood vessels was counted in five high-power fields using the “HOT SPOT” method. The hotspot areas in the tumor tissues are those that show the greatest microvessel density as defined by Weidner in 1991. Briefly, microvessel density was determined by light microscopy in areas of invasive tumor containing the highest numbers of capillaries and small venules (microvessels) per area (i.e., areas of the most intense vascularization). We found these vascular “hotspots” by scanning the tumor sections at low power (×40 and ×100) and identifying those areas of invasive carcinoma having the greatest numbers of distinct HIF staining microvessels per area [Figure 3]. The hotspots could occur anywhere within an invasive tumor but most frequently appeared at the margins of the carcinoma.[9] Then the mean was calculated as:

\[
\text{MVD} = \frac{\text{Total number of blood vessels counted}}{\text{Number of fields counted}}
\]

**Statistical analysis**

The correlation between MVD and HIF-1α staining was established using the Mann–Whitney test in all the three study groups. The intergroup correlation was also established.[10]

**RESULTS**

On comparing HIF-1α expression in OSMF, OSCC and OSCC with OSMF, a gradual increase was found in its intensity [Table 1].

The blood vessel density was found to be the highest in OSCC with OSMF, intermediate for OSCC and the lowest for OSMF [Table 2]. The HIF-1α expression and blood vessel density were positively correlated in all the three groups that are OSMF, OSCC and OSCC with OSMF [Tables 3].

**DISCUSSION**

OSMF is histopathologically characterized by fibrosis of subepithelial connective tissue. Collagens are the major structural component of extracellular matrix, hence precise regulation of collagen metabolism is essential to maintain the normal integrity of connective tissue.[5] With the progression of the disease process of OSMF, the production of collagen type 1 is increased and the degradation of collagen is reduced by up to 75%. Extensive fibrosis of the connective tissue causes reduction of vascularity, resulting in subsequent hypoxia in both fibroblasts and surface epithelium. Hypoxia causes atrophy and ulceration of the epithelium by inducing apoptosis. In addition, the overexpression of hypoxia-induced factor-1α is seen in OSMF, which indicates changes in cell proliferation, maturation and metabolic adaptation, increasing the possibility of malignant transformation.[11] The cellular response to hypoxic stress is controlled by a family of prolyl hydroxylases (PHD). In the presence of adequate oxygen, PHDs hydroxylate HIF-1α at conserved proline residues within the oxygen-dependent degradation domain. Once hydroxylated, HIF-1α becomes a substrate for von Hippel-Lindau–mediated ubiquitination and degradation. Upon degradation, HIF-1α...
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The hypoxic and anaerobic conditions found at the center of the solid tumor would require cells present there to rely on glycolysis. As the $O_2$ levels decreased, the generation of adenosine triphosphate (ATP) shifted from oxidative phosphorylation (OXPHOS) to glycolysis (Pasteur effect). OXPHOS is more efficient in generating ATP than glycolysis; the oxidation of one molecule of glucose gives a net yield of 30 ATPs via OXPHOS and 2 ATPs via glycolysis. It is comprehensible, therefore, that cells generate ATP through OXPHOS when they have enough $O_2$ levels.  

A hallmark of malignant tumors is the elevated uptake of glucose even under normal oxygen conditions, known as aerobic glycolysis or the “Warburg effect.” The Warburg effect is the metabolic change observed in cancer cells from OXPHOS to glycolysis as the primary source of cellular energy.  

It has been proposed that cancer cells have increased glycolytic rates despite the presence of $O_2$ because these cells have irreversible damages to OXPHOS. It has been reported that cells from the most common cancer types have decreased expression of ATP synthase, a protein complex required for OXPHOS. Also, mitochondrial mutations, which may lead to malfunction in OXPHOS, have been observed in tumor cells. It has also been shown that inactivation of p53 – one of the most commonly mutated genes in cancer – may trigger the Warburg effect; p53 is involved in the activity of cytochrome c oxidase, a protein complex involved in OXPHOS. These cellular responses have been shown to cause distinct transformations like the upregulation of proteins such as HIF-1$\alpha$ that help the tumor survive adverse conditions in which normal cells cannot persist.

Thus both hypoxic and normoxic cells in OSCC show a rise in HIF-1$\alpha$. The aerobic glycolysis stimulated by HIF-1$\alpha$ leads to a positive feedback loop in the cancer cells. Tumor cells in normoxic conditions show an over expression of HIF-1$\alpha$ when compared to normal cells in the same environment. The greater rate of aerobic glycolysis and a lower rate of oxidative

### Table 2: The mean blood vessel density found in various groups as defined by the HIF-1 staining intensity

| Group/criteria                  | Mild (patients) | Moderate (patients) | Severe (patients) |
|--------------------------------|-----------------|---------------------|-------------------|
| OSMF                           | 5-6 (02)        | 6-7 (10)            | 8-10 (08)         |
| OSCC                           | 7-8 (01)        | 10-11 (11)          | 14-15 (08)        |
| OSCC with OSMF                 | - (00)          | 12-13 (08)          | 15-17 (12)        |

HIF-1: Hypoxia inducible factor-1, OSMF: Oral submucous fibrosis, OSCC: Oral squamous cell carcinoma

### Table 3: Results of Mann-Whitney U-test

| Comparing between groups         | BvD       | Hpo       |
|----------------------------------|-----------|-----------|
| OSMF and OSCC                    |           |           |
| Mann-Whitney U-test              | 25.500    | 194.000   |
| Wilcoxon W                       | 235.500   | 404.000   |
| Z                                | −4.774    | −0.182    |
| Asymptotic significant (two-tailed)|       |           |
| Exact significant (2* (one-tailed significant)) | 0.000  | 0.883    |
| OSCC and OSCC with OSMF          |           |           |
| Mann-Whitney U-test              | 57.500    | 152.000   |
| Wilcoxon W                       | 267.500   | 362.000   |
| Z                                | −3.882    | −1.466    |
| Asymptotic significant (two-tailed)|       |           |
| Exact significant (2* (one-tailed significant)) | 0.000  | 0.143    |
| OSMF and OSCC with OSMF          |           |           |
| Mann-Whitney U-test              | 176.500   | 156.000   |
| Wilcoxon W                       | 386.500   | 366.000   |
| Z                                | −0.642    | −1.358    |
| Asymptotic significant (two-tailed)|       |           |
| Exact significant (2* (one-tailed significant)) | 0.521  | 0.242    |

* Significant value, OSMF: Oral submucous fibrosis, OSCC: Oral squamous cell carcinoma,
respiration in tumor cells leads to increased intracellular levels of glycolytic metabolites, specifically pyruvate. Pyruvate promotes the production and stability of HIF-1α thus creating a positive feedback loop that bolsters the proliferation of cancer cells.

Thus, the rise in HIF-1α expression in OSCC can be attributed to the genetic changes in the cancer cells along with the tumor hypoxia leading to stabilization and production of HIF-1α. We found that there was a gradual rise in the intensity of staining for HIF-1α among the three study groups. It was least for OSMF (mild-02, mod-10, sev-08 patients), intermediate for OSCC (mild-01, mod-11, sev-08 patients) and the highest for OSCC with OSMF (mod-08, sev-12 patients). This increased severity of HIF-1 expression in OSCC with OSMF can be due to the combined effect of these two pathways in OSCC with OSMF.

Angiogenesis is an important mediator of tumor progression. As tumors expand, diffusion distances from the existing vascular supply increases resulting in hypoxia. Sustained expansion of a tumor mass requires new blood vessel formation to provide rapidly proliferating tumor cells with an adequate supply of oxygen and metabolites, failure to do so deprive it of energy and nutrients. 

O2 acts as a molecular signal and through its availability, coordinates blood vessel growth with the metabolic demands of growing tumor mass. A large number of genes involved in different steps of angiogenesis have been shown to increase by hypoxia. The key regulator of hypoxia-induced angiogenesis is the transcription factor HIF-1. Multiple HIF-1 target genes such as vascular endothelial growth factor (VEGF), angiopoietin and fibroblast growth factor have been shown to modulate angiogenesis by promoting the mitogenic and migratory activities of endothelial cells. Angiogenesis has been shown to increase by hypoxia. The key regulator of hypoxia-induced angiogenesis is the transcription factor HIF-1α. Neo-angiogenesis is an essential step for many physiological processes such as growth, wound healing, organ regeneration and reproductive functions. Abnormal blood vessel growth occurs in several pathological conditions, including tumor growth and metastasis.

There was a gradual rise in the mean blood vessel density from OSMF to OSCC and a further rise was seen in OSCC with OSMF. This can be attributed to the similarly rising levels of HIF-1α. Tumor angiogenesis is a critical step in the development, metastatic spread and regrowth of cancer. Tumors promote angiogenesis by secreting factors, including VEGF-A, basic fibroblast growth factor and transforming growth factor-β. Several reports suggest that an angiogenic switch may be activated in the premalignant stage of several human cancers. Thus, tumor angiogenesis is not necessarily a characteristic of the invasive tumor, as originally thought, but may be an early event during cancer development.

CONCLUSION

A progressive increase in the HIF-1 and mean blood vessel density was observed in OSMF to OSCC. The combination of the two (OSCC with OSMF) lead to a tissue phenotype that shows more hypoxia and angiogenesis. Thus, we postulate that on conversion of premalignancy to malignancy it leads to a more aggressive tumor phenotype. Hypoxia appears to be a key factor in regulating this domino effect.

OSMF → HYPOXIA → OXYGEN THERAPY → PREVENTION OF OSCC

Hence, the treatment modalities addressing it would be more beneficial to help prevent this conversion.

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Conflicts of interest

There are no conflicts of interest.

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