Complete surgical excision of the tumor proper along with the removal of the surgical margins is the treatment of choice for oral squamous cell carcinoma (OSCC). However, the rate of local recurrence remains high (30%) even though histologically negative surgical margins (HNMs) have been achieved. Hence, molecular evaluation of surgical margins to detect residual cancer cells or a field of genetically altered cells is of paramount importance. Mutation in p53 gene is an early event in carcinogenesis that can precede clinically evident morphological changes. Thus, its detection in HNMs may indicate an impending malignancy. However, till date, literature is lacking regarding the immunoexpression of p53 in HNMs and its comparison with adjacent OSCC. Hence, the aim of the present study was to study p53 immunoexpression in OSCC and its HNMs. We included 12 paraffin-embedded tissue blocks of OSCC having tumor tissue and HNMs. The samples were subjected to immunohistochemical staining using primary mouse monoclonal antibody against p53, and the stained slides were evaluated for staining intensity and percentage of expression. Descriptive analysis and Chi-square test were applied. The expression of p53 was observed in 66% of HNMs and 91.6% of tumor tissue. The three cases which showed local recurrence demonstrated the expression of p53 in HNM. Hence, molecular analysis of p53 in surgical margins might potentially predict local recurrence of OSCC.

Keywords: Immunohistochemistry, oral squamous cell carcinoma, p53 protein, recurrence, surgical margins
of the Department of Oncopathology, Karnataka Cancer Therapy and Research Institute, Navanagar, Hubli, Karnataka. HNM was defined as the distance from invasive carcinoma to surgical margin of ≥5 mm. Cases affecting the gingivobuccal complex or buccal mucosa classified as T₁/T₂/T₃, N₁/N₂/N₃, M₁ were included. Margins showing evidence of epithelial dysplasia, patients who had received radiotherapy or chemotherapy prior to surgery, histological variants of OSCC and recurrent tumors were excluded. Five micron thick sections from each paraffin-embedded block were subjected to p53 immunostaining (DAKO). The immunohistochemistry stained slides were evaluated based on intensity of staining and number of positively stained cells. In tumor proper, p53 expression was considered positive if nuclear expression was noted in more than or equal to 20 tumor cells. For HNMs, brown staining of nuclei in more than 50 parabasal/suprabasal cells was considered positive. Varying degrees of p53 expression were noted in 11 out of 12 cases (91.6%) of tumor proper [Figure 1a], while one case showed no expression (8.3%). Comparison of p53 expression with clinicopathological parameters is depicted in Table 1. The correlation of p53 immunoexpression with age and habits is depicted in Table 2.

When HNMs were evaluated for immunoexpression, eight cases (66%) expressed p53, whereas four cases were negative for p53. Among the eight HNMs positive for p53, three cases showed expression only in basal layer [Figure 1b], three in both basal and parabasal layers [Figure 1c], and two cases demonstrated immunoreactivity in the entire thickness of epithelium [Figure 1d]. Interestingly, all three cases with recurrence/death demonstrated overexpression of p53 in HNM.

Most of the patients (9 out of 12), with p53 expression in tumor tissue and HNMs, were >40 years of age and practiced tobacco and alcohol consumption habits [Table 2]. This can be attributed to genetic modifications owing to the long term exposure of carcinogens such as polycyclic aromatic hydrocarbons.[1] However, the results were statistically insignificant, which might be due to a smaller sample size.

In our study, eight out of 12 cases (66%) showed p53 expression in HNMs. Studies by Singh et al.[8] (54.2%), Bilde et al.[3] (75%), Wang et al.[1] (31%) have also found p53 expression in HNMs. Five out of eight (62.5%) immunopositive HNMs showed basal and suprabasal p53 staining. Bilde et al.[3] reported similar results in their samples. Immunopositive cells in the tissue adjacent to malignant epithelium represent genetically stressed cells and are early indication of carcinogenesis; thus reflecting field canzerization.[2]

In the present study, p53 expression was observed in 91.6% of tumor tissue and 66% of HNMs. Except for one case, in all the immunopositive HNMs, the respective tumor tissue also showed p53 Immunoeexpression. Bilde et al.[3] reported that, of the 12 HNMs which were immunopositive for p53, seven HNMs had corresponding tumour tissue positive for p53. It has been hypothesized that p53 expressing cells are a part of preneoplastic field that escapes normal growth control gain growth advantage develops an expanding clone ultimately leading to recurrent tumor.[2]

Negative expression in adjacent tumor in some cases could be due to lack of representative tumor tissue sample, or due to the role of different cell cycle markers involved in the development of cancer. The presence of p53 immunoreaction only in HNM and not in the adjacent tumor, require further research.

However, in a study by Cruz et al.[6,7] all the cases showed p53 immunoeexpression in both HNMs and adjacent tumor. They also found that three of the HNMs with suprabasal p53 staining later developed carcinomas.[6,7] In the present study, recurrence in two patients and death in one patient was noted. Among these three cases, two showed immunoreaction in both HNMs and tumor tissue, and one
showed p53 expression only in HNMs. This indicates that analysis of molecular nature of the HNMs is of paramount importance to identify the genetically altered fields.

However, it is noteworthy that in spite of the high frequency of immunopositive HNMs, only a few patients developed recurrence. The possible reason could be due to the fact that a large number of patients undergo postoperative radiotherapy and chemotherapy, that eradicate residual cancer cells.\textsuperscript{13,14}

Overall, the results of the present study suggest that besides the gold standard histopathological assessment, the inclusion of “Molecular Status” of HNMs is of value to identify patients at risk of developing recurrence. Future prospective studies with a larger sample size with inclusion of additional margins may be needed to confirm our findings. In addition, combining TP 53 gene analysis by methods like PCR along with p53 protein in HNMs might improve the predictive value of p53 overexpression.

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Conflicts of interest
There are no conflicts of interest.

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