# Immunohistochemical expression of heparin-binding protein 17/ fibroblast growth factor-binding protein-1 (HBp17/FGFBP-1) as an angiogenic factor in head and neck tumorigenesis

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Abstract. Heparin-binding protein 17/fibroblast growth factorbinding protein-1 (HBp17/FGFBP-1) is a secreted protein that releases immobilized fibroblast growth factor-2 (FGF-2) from the extracellular matrix and plays a critical role in FGF bioactivation. In the present study co-localization of FGF-2 and HBp17/FGFBP-1 was observed in oral tissues including normal mucosa, hyperplasia, dysplasia of different degrees and oral squamous cell carcinoma (OSCC). The expression score for HBp17/FGFBP-1, FGF-2 as well as vascular endothelial growth factor A (VEGF-A) became higher with the severity of epithelial dysplasia and was highest in severe dysplasia. The expression of HBp17/FGFBP-1, FGF-2 and VEGF-A showed significant association with microvessel density, but no correlation with TNM stages or OSCC recurrence interval. Our results demonstrated that HBp17/ FGFBP-1, like VEGF-A and FGF-2, might also promote the induction of tumor angiogenesis. The strongest expression of angiogenic factors in severe dysplasia suggests a potential point for targeting novel anti-angiogenic therapeutic strategies.

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*Abbreviations*: HBp17, heparin-binding protein 17; FGF, fibroblast growth factor; FGFBP-1, fibroblast growth factor binding protein-1; VEGF-A, vascular endothelial growth factor A; OSCC, oral squamous cell carcinoma; MVD, microvessel density

*Key words:* premalignancy, angiogenesis, FGF-2, HBp17/ FGFBP-1, VEGF-A

# Introduction

Neovascularization is a critical event in tumor progression such that tumor growth beyond a few millimeters requires an adequate vasculature (1). FGF-2 and VEGF-A have been regarded as likely candidates for inducers of angiogenesis. VEGF-A is a positive regulator of tumor-associated angiogenesis, and it stimulates endothelial cell proliferation and enhances vascular permeability. Many reports have shown that VEGF-A is closely related to growth and prognosis in oral cancers (2-5). FGF-2 is the prototypic member of a family containing at least 23 structurally-related polypeptide growth factors (6). It is expressed ubiquitously in cells of mesoderm and neuroectodermal origin, and in a variety of tumor cells. In vitro, FGF-2 is a potent mitogen for different cell types, including vascular endothelial cells, fibroblasts and cancer cells such as squamous cell carcinoma (7) and salivary adenocarcinoma (8,9). In vivo, FGF-2 is a potent inducer of angiogenesis and has pleiotropic effects both on development, differentiation in various organs, and tumorigenesis (10,11).

FGF-2 is secreted despite lacking a leader sequence, which targets intracellular proteins for secretion to the extracellular environment, and externalized FGF-2 needs to be released and solubilized from the extracellular matrix. We initially isolated heparin-binding protein of 17 kDa in 1991 from the medium conditioned by human epidermoid carcinoma A431 cells and designated the protein as heparin-binding protein 17 (HBp17) (12). HBp17 was found to bind FGF-1 and FGF-2 in a non-covalent and reversible manner (12,13). Owing to its capacity to bind FGFs, HBp17 was re-named fibroblast growth factor-binding protein-1 (FGFBP-1) (14). Binding of HBp17/FGFBP-1 to FGF-2 mediates the release of immobilized FGF-2 from the extracellular matrix allowing it to reach its receptor, suggesting that HBp17/FGFBP-1 serves as an extracellular switch molecule for FGF-2 (12-17). Recently, it was reported that HBp17/FGFBP-1 also interacted with FGF-7, -10, and -22, and regulates FGF activities (18). Upregulation of HBp17/FGFBP-1 was found in various tumors, including head and neck, skin and colon cancers (19-21). In

Histopathological	Number	Average age	Gender		
grade	of cases	(years)	Male	Female	
Normal gingiva	10	24.3	3	7	
Hyperplasia	20	58.7	10	10	
Moderate dysplasia	20	65.8	12	8	
Severe dysplasia	14	61.7	8	6	
OSCC	30	64.2	20	10	

Table I. Patient characteristics.

addition, the growth and angiogenesis of xenografted tumors in mice was decreased in parallel with a reduction in HBp17/ FGFBP-1 expression, suggesting that human tumors utilize HBp17/FGFBP-1 as an angiogenic switch molecule (22).

Although angiogenesis has been shown to be an independent prognostic indicator in a number of cancers, little is known about its significance in premalignant lesions in the head and neck region. To our knowledge there have been no reports on the role of HBp17/FGFBP-1 in premalignant lesions. The aim of this study was to investigate the expr-ession of HBp17/ FGFBP-1, FGF-2 and VEGF-A in the multistep process of head and neck tumorigenesis, and to evaluate the relationships between the expression of these angiogenic factors and tumor microvessel densities, TNM stages, OSCC recurrence interval and clinicopathological stages.

#### Materials and methods

Selection of specimens. Paraffin-embedded specimens representing premalignant lesions and oral squamous cell carcinoma (OSCC) were prepared in the Center for Oral Clinical Examination, Hiroshima University Hospital after surgical excision in the Department of Molecular Oral Medicine and Maxillofacial Surgery at Hiroshima University Dental Hospital. Tissue samples of normal oral epithelium were obtained from surgical specimens of non-cancer patients. The Ethics Committee in Hiroshima University approved specimen collection. All tissue samples were fixed in 10% neutral formalin for 24 h. Sections were prepared at 4  $\mu$ m in thickness for immunohistochemical procedures.

Antibodies. A monoclonal anti-HBp17 antibody designated as C9 (10) was used in this study. C9 hybridoma cells were cultured in RD medium [RPMI-1640 medium: Dulbecco's modified Eagle's medium (DMEM); 1:1] supplemented with 10  $\mu$ g/ml insulin, 5  $\mu$ g/ml transferrin, 10  $\mu$  2-mercaptoethanol, 10  $\mu$ M 2-aminoethanol and 10 nM selenite (RD+5F). After being filtered with a 0.45- $\mu$ m filter (Millipore), the conditioned medium was passed through Protein G affinity chromatography (Amersham Bioscience, Buckinghamshire, UK), and HBp17 antibody C9 was eluted by 0.1 M glycine- HCl (pH 2.7). After being purified, 10  $\mu$ g/ml of the C9 antibody was used for immunostaining.

Monoclonal anti-FGF-2 antibody (clone bFM-1, diluted 1:100; final concentration, 2  $\mu$ g/ml) was purchased from Upstate Biotechnology Inc. (Lake Placid, NY). Monoclonal anti-VEGF-A antibody (clone R-11, diluted 1:100; final

concentration, 2  $\mu$ g/ml) was from IBL (Int. Gunma, Japan). Monoclonal anti-von Willebrand factor antibody (vWF, diluted 1:40; final concentration, 5.75  $\mu$ g/ml) was purchased from Dako (A/S, Denmark).

Immunohistochemistry. The expression of HBp17/FGFBP-1, FGF-2, VEGF-A and vWF were analyzed with a three-stage avidin-biotin immunoperoxidase technique using a vectastain mouse ABC kit (Vector Laboratories, Burlingame, CA, USA). Briefly, deparaffinized tissue sections were incubated with 1% H<sub>2</sub>O<sub>2</sub> in methanol for 30 min at room temperature to block endogenous peroxidase activity, followed by incubation with 1.5% normal horse serum at 37°C for 30 min to prevent non-specific binding. Incubation with primary antibody was performed at 37°C for 60 min using predetermined optimal dilution. As negative controls, sections were probed with 1.5% normal horse serum. Sections were then incubated with biotinylated anti-mouse immunoglobulin for 60 min, followed by 30-min incubation with Avidin-Biotin-peroxidase-Complex at room temperature. Reaction products were visualized by immersing sections in 10 mM Tris-HCl (pH.7.4) containing 1 mg/ml 3,3 diaminobenzidine (DAB) and 0.03% hydrogen peroxide. The sections were counterstained with Mayer's solution.

*Expression score*. A semi-quantitative method was used to evaluate the degree of immunostaining ranging from - (no expression), + (weak), ++ (moderate), to +++ (highest level of expression) on a cell-by-cell basis in 5 microscopic fields (x33 magnification). For each section, the percentage of cells stained at x33 magnification for five different fields were multiplied by that intensity value and the five products determined were added together to obtain a weighed average staining score.

*Microvessel density (MVD).* The microvessels were highlighted by staining endothelial cells with anti-vWF antibody by immunohistochemistry. MVD were counted at x200 magnification for ten fields and the mean value of those ten fields in each sample was calculated.

Statistical analysis. Data are presented as mean  $\pm$  SD. The relationship between FGF2, HBp17/FGFBP-1, VEGF-A expression and MVD in cancer and premalignant lesions was determined with the Spearman correlation test. The Kruskal-Wallis test was used when comparing immunohistological expression in the TNM stages of oral squamous cell carcinoma groups. All tests were three-tailed, and P≤0.05 was considered to indicate statistical significance.

## Results

Thirty specimens of OSCC, 54 specimens of premalignant oral lesions including 20 hyperplasias, 20 moderate dysplasias, 14 severe dysplasias and 10 control specimens were studied. There were 53 male and 41 female subjects, and the average age of the patients was 54.9 years old (Table I).

HBp17/FGFBP-1 expression. Fifty percent of normal gingiva showed weak expression at the cytoplasm of the spinous layer

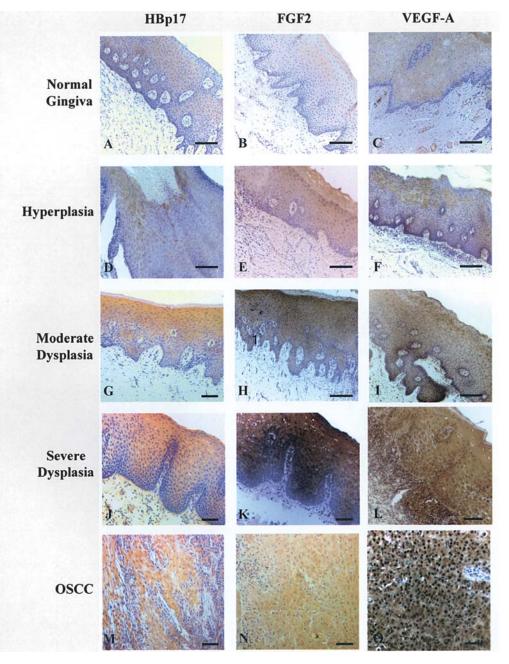


Figure 1. Immunohistochemical expression of HBp17/FGFBP-1, FGF-2 and VEGF-A in normal gingivas, hyperplasias, moderate dysplasias, severe dysplasias and oral squamous cell carcinomas. Normal gingiva showed weak expression of HBp17/FGFBP-1, FGF-2 and VEGF-A in the cytoplasm of spinous layer cells (Fig. 1A, B and C). Hyperplasias showed limited expression of HBp17/FGFBP-1, FGF-2 and VEGF-A in the acanthotic epithelium (Fig. 1D, E and F). Moderate dysplasias, severe dysplasias and OSCC strongly expressed HBp17/FGFBP-1 and FGF-2 in the cytoplasm of cells (Fig. 1G, H, J, K, M and N). VEGF-A expression was detected not only in the cytoplasm but also in cell nuclei in moderate dysplasia, severe dysplasia and OSCC tumors (Fig. 1I, L and O). Scale bar, 50  $\mu$ m.

Table II. Immunostaining intensities for HBp17/FGFBP-1, FGF-2 and VEGF-A in normal gingivas, hyperplasias, moderate dysplasias, severe dysplasias and oral squamous cell carcinomas.

	HBpl7						FGF-2				VEGFA				
Cases	-	+	++	+++	(% positive)	-	+	++	+++	(% positive)	-	+	++	+++	(% positive)
Normal gingiva	5	5	0	0	(50)	6	4	0	0	(40)	5	5	0	0	(50)
Hyperplasia	8	6	6	0	(60)	12	6	2	0	(40)	10	10	0	0	(50)
Moderate dysplasia	6	8	6	0	(70)	10	4	4	2	(50)	8	4	6	2	(60)
Severe dysplasia	0	0	4	10	(100)	0	0	8	6	(100)	2	2	4	6	(85.7)
OSCC	0	0	22	8	(100)	0	2	22	6	(100)	6	2	14	8	(80)

\* \*

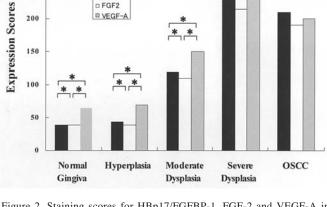


Figure 2. Staining scores for HBp17/FGFBP-1, FGF-2 and VEGF-A in multisteps of oral tumorigenesis. Staining scores gradually increased as gingival lesions proceeded from normal to hyperplasia, moderate dysplasia, severe dysplasia and OSCC. Staining scores were highest in severe dysplasia. At all stages of oral tumorigenesis, staining scores of HBp17/ FGFBP-1 and FGF-2 showed significant correlations.

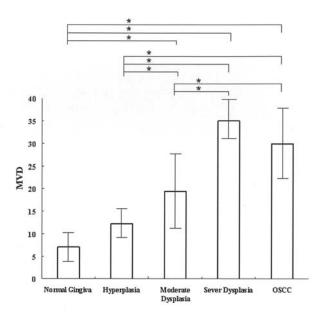


Figure 3. Microvessel density (MVD) in multisteps of oral tumorigenesis. MVD increased gradually with the severity of epithelial dysplasia, \*p<0.05.

(Fig. 1A). Sixty percent of hyperplasias showed limited expression at the acanthotic epithelium (Fig. 1D). Seventy percent of moderate dysplasias and 100% of severe dysplasias and OSCC strongly expressed HBp17/FGFBP-1 in the cytoplasm (Fig. 1G, J, M and Table II).

FGF-2 expression. Forty percent of normal tissue expressed FGF-2 in the cytoplasm in the spinous layer (Fig. 1B) and hyperplasias expressed FGF-2 in the cytoplasm in the acanthotic epithelium (Fig. 1E). The staining intensity was greater in the upper layers of moderate dysplasias (Fig. 1H), and FGF-2 staining significantly increased in intensity through to the severe dysplasia (Fig. 1K). In OSCC, FGF-2 expression was detected in the cytoplasm of tumor cells (Fig. 1N). These Table III. Relationships between MVD and expression of HBp17/FGFBP-1, FGF-2 and VEGF-A at all stages of tumorigenesis.<sup>a</sup>

	Expression score	MVD (mean±SD)	P-value
Hyperplasia			
HBpl7	45		
FGF-2	40	12.2±3.56	*
VEGF-A	70		
Moderate dysplasia			
HBpl7	120		
FGF-2	110	19.4±8.9	*
VEGF-A	150		
Severe dysplasia			
HBpl7	230		
FGF-2	215	35.1±4.8	*
VEGF-A	240		
OSCC			
HBpl7	210		
FGF-2	190	30.1±7.3	*
VEGF-A	200		

<sup>a</sup>Expression scores are mean values, \*p<0.05.

data suggested that FGF-2 co-localized with HBp17/FGFBP-1 in both normal and tumor tissues.

VEGF-A expression. The immunohistochemical staining of VEGF-A appeared to be localized to the cytoplasm of normal control cells, hyperplasias, and endothelial cells lining microvessels (Fig. 1C and F). In the case of dysplasias of different degrees, the level and extent of VEGF-A expression increased with the severity of epithelial dysplasia. VEGF-A expression was detected not only in the cytoplasm but also in the nuclei of tumor cells (Fig. 1I, L and O). As the grade of dysplasia increased, the extent and levels of expression of HBp17/ FGFBP-1, FGF2 and VEGF-A also increased. The expression of HBp17/FGFBP-1, FGF-2 and VEGF-A showed significant correlations (P<0.05) at all stages (Fig. 2).

Microvessel density (MVD) measured by vWF expression gradually increased with the severity of epithelial dysplasia. MVD was  $7.02\pm3.04$  in normal gingival,  $12.2\pm3.56$  in hyperplasias, 19.4±8.9 in moderate dysplasias, 35.1±4.8 in severe dysplasias, and 30.1±7.3 in OSCC. MVD was the highest in severe dysplasias (Fig. 3). The expression of HBp17/FGFBP-1, FGF-2 and VEGF-A showed significant correlation with MVD at all steps of tumorigenesis (Table III). In contrast, no associations were found between expression score and TNM stages or recurrence interval (data not shown).

## Discussion

We have previously reported that limited expression of FGF-1 and FGF-2 was detected in the spinous layer of normal gingiva,

300

250

200

HBp17

G FGF2 VEGE-A and stronger expression was detected in the cytoplasm of oral squamous cell carcinomas by immunohistochemistry (7). A second study also suggested that HBp17/FGFBP-1 mRNA was detected not only in tumor cells but also in keratinocytes of normal mucosa (23). The results of the present study agree with the earlier findings in demonstrating that co-localization of FGF-2 and HBp17/FGFBP-1 was observed in the spinous layer of normal gingiva. The extent and intensity of expression of these two markers increased with the severity of epithelial dysplasia. Therefore, we propose that HBp17/FGFBP-1 in collaboration with FGF-2 has a regulatory role in tumorigenesis and normal development in epithelial tissue.

In a variety of cancers angiogenic factors have become significant prognostic markers (24-28). Our study showed that the expression of the angiogenic molecules HBp17/FGFBP-1, FGF-2 and VEGF-A also increased with the severity of epithelial dysplasia, and the expression score was highly correlated at all stages of the multistep development of head and neck cancer. The expression score was highest in severe dysplasia, but not in OSCC. This difference may be due to accumulated genetic alterations within tumor cells leading to growth factor independence. Andreas *et al* also reported an increase in HBp17/FGFBP-1 expression during development and tumorigenesis and a relative decrease in HBp17/FGFBP-1 expression in later stages of carcinogenesis (29).

For evaluating angiogenesis during neoplastic development, we measured MVD and used it to represent the angiogenic status of lesions. We found a significant association between MVD and expression of HBp17/FGFBP-1, FGF-2, and VEGF-A. These data are supported by a study of head and neck cancer in which Li *et al* reported a significant association between increased expression of VEGF or FGF-2 and MVD in different histological stages of head and neck tumorigenesis (30). Frank *et al* reported that enhanced expression of FGF-2 and FGF-2 receptors by cancer cells and vascular endothelial cells promoted cancer growth and metastasis through angiogenesis (31). Subsequently, Fukuiwa *et al* found a significant correlation between VEGF expression and MVD (32).

Our results showed no association between angiogenic factor expression and recurrence intervals or TNM stages (data not shown). Another study found that the elevated level of both IL-8 and VEGF correlated with an increased incidence of recurrence, shorter disease-free intervals and higher initial TNM stage in head and neck tumorigenesis (33). However, Petri et al found no association with VEGF expression and TNM stages or MVD (34). Similarly, Burian et al observed no association between FGF-2 expression and TNM stages or MVD (35). In the present study, a significant increase in MVD observed in severe dysplasia rather than in moderate dysplasia and in OSCC. The decrease in MVD in tumors compared to severe dysplasia may be due to a decreased requirement for further angiogenesis once a certain stage in the disease process has been reached. This idea is supported by Li et al who also found increased expression of MVD in severe dysplasia compared to tumor specimens (30).

In conclusion, our results suggest that the angiogenic factors HBp17/FGFBP-1, FGF-2 and VEGF-A may together provide essential regulatory signals for the induction of tumor angiogenesis. The strongest expression of angiogenic factors and

highest MVD in the severe dysplasia stage suggest potential targets for therapeutic strategies. Thus, the appropriate choice of therapy before or at the stage of severe dysplasia might prevent further malignant progression by controlling angiogenesis.

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