

Orchestrated expression of vasculogenic mimicry and laminin-5 γ 2 is an independent prognostic marker in oral squamous cell carcinoma

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Abstract

Vasculogenic mimicry (VM), an endothelial cell-independent alternative mechanism of blood supply to the malignant tumour, has long been considered as an adverse prognostic factor in many cancers. The correlation of VM with laminin-5 γ 2 and the assessment of their harmonized expression as an independent risk factor have not been elucidated yet in oral squamous cell carcinoma (OSCC). CD31/PAS staining stratified 116 clinically diagnosed OSCC specimens into VM+ and VM- cohorts. The expression pattern of laminin-5 γ 2 and its upstream modulator MMP2 was evaluated by immunohistochemistry and Western blot. The Kaplan-Meier and Cox regression analyses were performed to assess the survival and prognostic implications. The presence of VM demonstrated a significant correlation with the expression of laminin-5 γ 2 ($p < .001$) and MMP2 ($p < .001$). This pattern was mirrored by the significant upregulation of laminin-5 γ 2 and MMP2 in VM+ cohorts compared with the VM- ones. Furthermore, co-expression of VM and laminin-5 γ 2 was significantly associated with tumour grade ($p = .010$), primary tumour size ($p < .001$), lymph node metastasis ($p = .001$) and TNM stages ($p < .001$) but not with patients' age, gender, tobacco and alcohol consumption habit. Vasculogenic mimicry and laminin-5 γ 2 double-positive cohort displayed a significantly poorer disease-free survival (DFS) and overall survival (OS). Vasculogenic mimicry, laminin-5 γ 2 and their subsequent dual expression underlie a significant prognostic value for DFS [hazard ratio (HR) = 9.896, $p = .028$] and OS [HR = 21.401, $p = .033$] in OSCC patients. Together, our findings imply that VM along with laminin-5 γ 2 is strongly linked to the malignant progression in OSCC and VM and laminin-5 γ 2 coordination emerges as a critical prognostic biomarker for OSCC.

KEYWORDS

co-expression, laminin-5 γ 2, OSCC, prognosis, survival, vasculogenic mimicry

1 | INTRODUCTION

Oral cancer is the 6th most common malignancy in the world.¹ Oral squamous cell carcinoma (OSCC) accounting for over 90% of oral cancers² is one of the most common causes of cancer-related deaths in the developing countries including India and the South-East Asia. The estimated incidence and mortality due to lip, oral cavity cancer in the world are 2.0% and 1.8% respectively.³ Annually 75,000–80,000 new oral cancer cases are being reported in India.⁴ The use of tobacco in various forms including cigarette, bidi, hookah, betel nut and betel quid is the major risk factor for OSCC.⁵ Severe alcoholism, HPV infection, dietary deficiencies and poor oral hygiene are the other common identified risk factors.⁶ Metastasis and postoperative recurrence are the most common reasons for poor 5-year survival⁷ that further increases failure of treatment. There are multiple clinicopathological factors responsible for this poor outcome. Growing tumours survive in the nutrient- and oxygen-deficit state using diverse strategies. In 1999, Maniotis et al.⁸ demonstrated that when blood supply cannot meet the need of rapid tumour growth, some aggressive, metastatic and genetically dysregulated cancer cells mimic the endothelial cells and form pseudovascular channel-like structures called vasculogenic mimicry (VM). It was first described in human uveal melanoma as periodic acid–Schiff (PAS)–positive patterned vascular network and enables the tumours to form matrix-embedded vascular structures carrying plasma and blood cells to fulfil the increasing nutrient and metabolic demands in tumour microenvironment.⁹ Core matrix proteins such as laminin, heparan sulphate proteoglycan and collagens IV and VI have been identified in these patterns.¹⁰ Though vasculogenic mimicry is considered as an established prognostic marker in melanoma,^{11,12} breast cancer,^{13,14} glioblastoma,¹⁵ ovarian cancer,¹⁶ lung cancer,¹⁷ prostate cancer¹⁸ and digestive cancers,¹⁹ the underlying molecular phenotypes inducing it in OSCC and their prognostic significance are poorly understood.

Laminin-5 (Ln-5), a component of extracellular matrix (ECM) adhesion molecules, is expressed predominantly in the epithelial basal membrane structure that promotes static adhesion and hemidesmosome formation.²⁰ However, the cleavage of short arm of $\gamma 2$ subunit of laminin-5 by matrix metalloproteinases (MMPs) such as MMP2 and MT1-MMP leads to its switching from static to migratory form resulting in cell migration and/or invasion.^{21,22} In the context of the molecular mechanism influencing VM, the downstream signalling of VE-cadherin, EphA2 and VEGF choreographs the cleavage of laminin-5 into the pro-migratory $\gamma 2x$ and $\gamma 2'$ fragments through

activated MMP2, implicating the roles of the extracellular matrix remodelling in positively regulating the formation of VM network.²³

Although the differential expression of laminin-5 $\gamma 2$ has been associated with tumour invasion and lymph node metastasis in OSCC^{24,25} and with the poor survival outcome in TCGA database–derived head and neck cancer (HNC) cohorts ($n = 502$),²⁶ its correlation with VM phenotype and the prognostic significance of their coordinated expression have not been elucidated yet. Here, we aim to investigate the phenotypic characteristics of VM structures in OSCC tumour tissues and evaluate the expression of laminin-5 $\gamma 2$ and its upstream modulator MMP2, as well as their correlation in the process of the acquisition of VM structure in OSCC microenvironment. Finally, we have undertaken the survival and risk factor assessment of VM-laminin-5 $\gamma 2$ coordination in a defined patient cohort to enlighten a novel and promising therapeutic target of OSCC.

2 | MATERIALS AND METHODS

2.1 | Patients and tissue samples

The surgical and clinically confirmed OSCC tissue specimens from a sum total of 116 patients (median age: 54 years, range: 28–80 years) were obtained from Chittaranjan National Cancer Institute, Kolkata, during May 2014–April, 2015. Patients with history of recurrence or preoperative chemotherapy and radiotherapy were excluded. Informed written consent was obtained from all the patients prior to specimen collection. The study was approved by the Institutional Ethics committee (IEC Ref: A-4.311/53/2014) in accordance with the ethical guidelines of Declaration of Helsinki (1964) and its later amendments. Tumour-node-metastasis stages were evaluated according to 8th edition of the American Joint Committee on Cancer (AJCC), and tumour grade was classified according to World Health Organization (WHO) standards. For OSCC patients with complete clinicopathological information and follow-up data, the overall survival (OS) time was calculated as the time interval from the date of surgery of the patients to their oral cancer–related death, and disease-free survival (DFS) was noted as the time interval from the date of surgery to the first documentation of local recurrences or distant metastasis. Parameters that are associated with diagnosis, prognosis and treatment of OSCC such as age, anatomic location of primary tumour, histological grade, habit of tobacco and alcohol consumption, tumour size, lymph node metastasis, and TNM stage group have been recorded (Table 1).

2.2 | Immunohistochemistry/PAS dual staining

Immunohistochemistry (IHC) coupled with PAS staining was performed with the primary antibodies against CD31 or PECAM-1 (Santa Cruz Biotechnology, Inc; rabbit monoclonal; clone: M-20, dilution: 1:100), laminin-5 (Y2 chain) (Merck; mouse monoclonal; clone: D4B5, dilution: 1:100) and MMP2 (Novus Biologicals; mouse monoclonal; clone: 8B4, dilution: 1:100) as per previously described methods.^{14,27}

2.3 | Evaluation of vasculogenic mimicry and IHC markers

Vasculogenic mimicry (VM) was identified through the detection of CD31-negative and PAS-positive lumen-like structures surrounded by tumour cells (but not with endothelial cells) with or without red blood cells inside the lumen²⁸⁻³² (Figure 1A). The vascular structures were observed for structural integrity with no incidence of haemorrhage, necrosis or inflammatory cell infiltration in close proximity.³³⁻³⁶ The VM density with respect to the overall vascular density has been assessed according to the modified method described by Weinder et al,³⁷ 1991, Shao et al, 2008,³⁸ and Zhou et al 2019.³⁹ The total number of CD31+ and CD31- lumen-like vascular structures, surrounded by tumour cells or endothelial cells, was considered as the overall vascular density. The areas of highest vascular density were found by observing the slides at 200× magnification. VM vessels were individually counted in 5 randomly selected 200× magnification field. The average percentage of VM has been evaluated relative to the overall vascular density and graded on the basis of following score: 0, negative; 1, <20%; 2, 20-<40%; 3, 40-<60%; and 4, ≥60%. The immunohistochemical score of our studied markers (laminin-5γ2 and MMP2) was determined by considering intensity of staining and proportion (%) of stained cells.⁴⁰⁻⁴⁵ All the staining results were blindly evaluated by two experienced pathologists in a semi-quantitative manner. To account the intra-tumoral heterogeneity of antibody expression, ten randomly selected represented fields (under 400× magnification) from different areas of each slide were evaluated by two qualified pathologists (manual method). The staining intensity was determined on the basis of the following score: 0, negative; 1, mild; 2, moderate; and 3, strong staining, and the percentage (proportion) of positively stained cells per field was scored as follows: 0, <10%; 1, <25% of positively stained cells; 2, <50% of positively stained cells; and 3, >50% of positive cells.⁴⁶⁻⁴⁸ The final immunohistochemical staining score of each sample was determined by summation of staining intensity and percentage (proportion) of positively stained cells, which ranged from 0 to 6. The final staining score 0-3 was considered as negative staining

TABLE 1 Demographic and clinicopathological profile of OSCC patients

Patients' characteristics	n (%)
Age (years)	
<55	62 (53.45)
≥55	54 (46.55)
Gender	
Male	85 (73.28)
Female	31 (26.72)
Tobacco consumption	
Yes	50 (43.10)
No	66 (56.90)
Alcohol consumption	
Yes	10 (8.62)
No	106 (91.38)
Tumour location	
Lip	8 (6.89)
Tongue	17 (14.65)
Buccal mucosa	39 (33.62)
Gingiva	24 (20.69)
Floor of mouth	4 (3.45)
Retromolar trigone (RMT)	5 (4.31)
Others ^a	19 (16.38)
Grade	
Well	73 (62.93)
Moderate	41 (35.34)
Poor	2 (1.72)
Primary tumour status	
T1	64 (55.17)
T2	8 (6.89)
T3	18 (15.52)
T4	26 (22.41)
Lymph node metastasis	
N0	66 (56.89)
N1	27 (23.27)
N2	21 (18.10)
N3	2 (1.72)
TNM stage group	
I (T1N0M0)	50 (43.10)
II (T2N0M0)	5 (4.31)
III (T3N0M0, T1-3N1M0)	23 (19.83)
IV (T4N0M0-T1-4N1-3M0)	38 (32.76)

^aOthers include alveolar mucosa, hard palate and soft palate.

and that of 4-6 was considered as positive staining.^{38,49} The semi-quantitative evaluation of pathologists was further validated by IHC profiler plugin⁵⁰ compatible with ImageJ (Figure S1). To normalize the digital image analysis with the

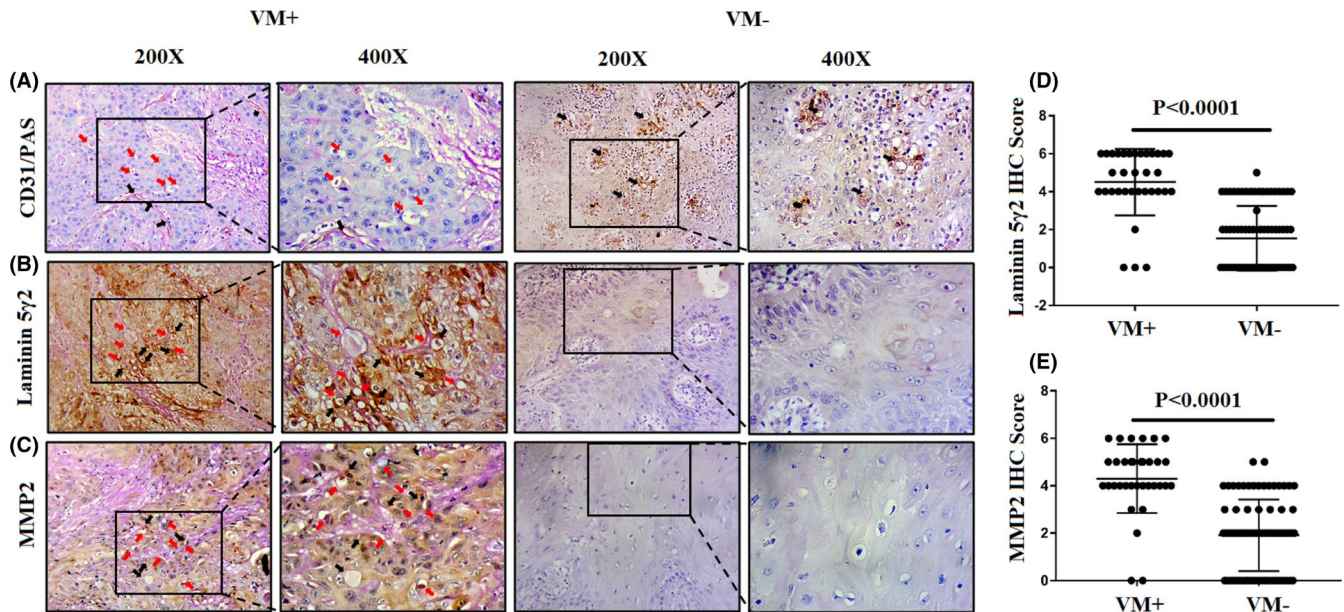


FIGURE 1 (A) CD31-PAS staining showing VM and endothelial structure in OSCC. Red arrows indicate PAS-positive and CD31-negative VM architecture, and black arrows represent endothelial structure showing CD31-positive staining with or without PAS staining (200× and 400× magnification). Representative images for immunohistochemical status (coupled with PAS staining) of (B) laminin-5γ2 and (C) MMP2 in VM-positive and VM-negative OSCC cohorts (200× and 400× magnification). Red arrows indicate PAS-positive networks, and black arrows indicate specific expression of proteins (B,C). Quantitative analysis of (D) laminin-5γ2 and (E) MMP2 expressions was revealed by immunohistochemical scores. The differences between VM-positive and VM-negative groups were calculated by the *t* test ($p < .0001$)

pathologists' manual analysis, the IHC profiler-generated four-tier staining pattern was scored from 0, negative; 1, low positive; 2, moderately positive; and 3, high positive similar to the manual assessment. Further, the total percentage of positive staining (1, low positive; 2, moderately positive; and 3, high positive) was determined from their individual percentage contribution and also scored from 0 to 3 scale in accordance with the pathologists' consideration of score for proportion (%) of stained cells.^{46–48} The final immunohistochemical score (0–6) of each sample was determined by summation of the percentage score of total positive staining and the total intensity score contributed by different degrees of positive staining pattern to harmonize the digital assessment with the manual findings inferring the final score ≤ 3 as negative staining and that of > 3 as positive staining.

2.4 | Western blot

Western blot analysis was performed using 3–5 mg of tissue specimens from few representative samples of both VM+ ($n = 15$) and VM- ($n = 15$) groups as per our previously described methods.⁵¹ Fifty micrograms of total protein extract and appropriate primary antibodies against laminin-5 (Y2 chain) (Merck; mouse monoclonal; Clone: D4B5, Dilution: 1:100) and MMP2 (Novus Biologicals; mouse monoclonal; Clone: 8B4, Dilution: 1:100) was used. β -Actin was used as a loading control.

2.5 | Statistical analysis

All the statistical analyses were performed using SPSS 17 software (SPSS Inc) and GraphPad Prism version 7.00 software. The chi-squared (χ^2) test was performed to find the associations between clinical-pathological parameters and VM and laminin-5γ2, and correlation among laminin-5γ2, MMP2 expression and VM was determined by the Spearman correlation test. The *t* test was used to compare two means. Kappa value was calculated to assess the agreement between two pathologists and two methods. The Kaplan–Meier survival analysis followed by the log rank test was used to compare the survival patterns. The multivariate Cox proportional hazard regression model was also used where overall survival (OS) and disease-free survival (DFS) were calculated. $p < .05$ was taken to be statistically significant.

3 | RESULTS

3.1 | Evaluation of VM in OSCC tissue specimens

Vasculogenic mimicry (VM) was identified in 29.31% of OSCC tissue specimens. Based on the CD31/PAS staining, the total patient population was stratified into VM-positive (VM+) and VM-negative (VM-) cohorts (Figure 1A). 34

of 116 (29.31%) cases were VM-positive, and 82 (70.69%) cases were VM-negative.

3.2 | Correlation of VM with the differential expression of laminin-5 γ 2 and MMP2

The immunohistochemical expression of laminin-5 γ 2 and MMP2 was observed in OSCC tumour cell cytoplasm and in tumour stroma. The expression of laminin-5 γ 2 and MMP2 was positively correlated to VM at their individual level ($r = .6642$, $p \leq .001$ and $r = .6201$, $p \leq .001$ respectively), and the expression of laminin-5 γ 2 and MMP2 was also significantly correlated ($r = .5046$ and $p \leq .001$) with each other (Figure S2A–C). The expression pattern of laminin-5 γ 2 and MMP2 in VM-positive and VM-negative cohorts has been represented in Figure 1B,C. The quantitative data indicated the significantly elevated expression of laminin-5 γ 2 and MMP2 in VM-positive cohorts ($p < .0001$; Figure 1D,E). The mean immunohistochemical score of laminin-5 γ 2 and MMP2 in VM-positive cohorts was 4.50 ± 1.75 and 4.29 ± 1.45 (mean \pm SD), respectively, whereas the same for the VM-negative cohorts was 1.74 ± 1.54 and 1.91 ± 1.51 respectively. Figure S1 shows the significantly positive correlation ($R^2 > .9$, $p < .001$) between the IHC scoring of two pathologists and between digital and manual methods. Kappa statistics was used to assess the interobserver and intermethod agreement, which revealed almost perfect strength of agreement ($\kappa = 0.81$ – 1.00) between the scoring of two pathologists and substantial strength of agreement ($\kappa = 0.61$ – 0.80)⁵² between digital and manual methods confirming the reliability of our IHC scoring method. The differential expressional status of the studied markers was also validated through quantitative Western blot analysis (Figure S2D,E), which indicated significantly higher intensity of immunoreactive bands in VM-positive groups compared with the VM-negative ones ($p < .05$).

3.3 | Association of VM and laminin-5 γ 2 with the clinicopathological features

The association of individual and coordinated expression of VM and laminin-5 γ 2 with the clinicopathological characteristics of patients has been summarized in Table 2. The result showed 30 (25.86%) VM–laminin-5 γ 2 double-positive cases, 4 (3.44%) VM-positive laminin-5 γ 2 negative cases, 21 (18.10%) VM-negative laminin-5 γ 2–positive cases and 61 (52.58%) VM–laminin-5 γ 2 double-negative cases. The presence of vasculogenic mimicry in OSCC was significantly associated with tumour grade ($p = .002$),

primary tumour status ($p < .001$), lymph node metastasis ($p = .005$) and TNM stage group ($p < .001$) but not with patient's age, sex and tobacco or alcohol consumption habit. It has been significantly found that 50% (1/2) cohort of the poorly differentiated tumour grade developed VM, whereas 48.78% (20/41) of the moderately differentiated group and 17.80% (13/73) of the well-differentiated group were found to be VM-positive. It is also noteworthy that 59.09% (26/44) patients of T3 and T4 primary tumour status significantly developed VM compared with the T1 and T2 group [11.11% (8/72)]. Similarly, the occurrence of VM was also significantly prevalent in the patients with positive nodal status compared with the negative ones [46% (23/50) vs 16.66% (11/66)], as well as in the patients with TNM stage groups III and IV [47.54% (29/61)] compared with TNM stage groups I and II [9.09% (5/55)]. The positive rate of VM was also significantly associated with the expression of laminin-5 γ 2 in OSCC. 43.9% of the total patients (51/116) showed positive expression of laminin-5 γ 2. Among them, the VM-positive group was found to have significantly increased level of expression of laminin-5 γ 2 [88.23% (30/34)] compared with the VM-negative [25.61% (21/82)] counterparts. Similar to the findings for the association of VM with the clinicopathological features, the positive expression of laminin-5 γ 2 was mostly observed in the patients with T3 and T4 primary tumour status (68.18%, $p < .001$) and TNM stage groups III and IV (57.37%, $p = .002$). Interestingly, the double-positive expressional status of VM–laminin-5 γ 2 was also significantly associated with tumour grade ($p = .010$), primary tumour status ($p < .001$), lymph node metastasis ($p = .00133$) and TNM stage group ($p < .001$) indicating the strong correlation between VM and laminin-5 γ 2 in the pathogenesis of OSCC. In addition to laminin-5 γ 2, we have also found a significant positive association of its upstream modulator MMP2 with VM ($p < .001$; Table S1). In association with the significant positive correlation of MMP2 with VM and laminin-5 γ 2, we have also demonstrated the significant association of MMP2 and VM–MMP2 dual positivity with the other established prognostic features of OSCC (Table S1). These data indicated the deterministic role of laminin-5 γ 2 and its activator molecule MMP2 in the occurrence of VM and progression of OSCC.

3.4 | Correlation of the positive expression of laminin-5 γ 2 and VM with disease-free and overall survival

To understand the collaborative prognostic significance of VM–laminin-5 γ 2, the 5-year survival rate was calculated for total of 116 patients with respect to DFS and OS. After completion of the follow-up (median follow-up

TABLE 2 Association between VM, laminin-5 γ 2 and their dual expression with clinicopathological characteristics of oral squamous cell carcinoma (OSCC)

Patients' characteristics	VM				Laminin-5 γ 2				VM and laminin-5 γ 2 dual expression					
	Positive		Negative		Positive		Negative		Positive		Negative		χ^2	p value
	n (%)	n (%)	n (%)	n (%)	n (%)	n (%)	n (%)	n (%)	n (%)	n (%)				
Age (years)														
<55	17 (14.65)	45 (38.79)	0.230	0.632	23 (19.83)	39 (33.62)	2.55	0.110	14 (12.07)	48 (41.38)	0.748	0.387		
\geq 55	17 (14.65)	37 (31.89)			28 (24.14)	26 (22.41)			16 (13.79)	38 (32.76)				
Gender														
Male	23 (19.83)	62 (53.45)	0.778	0.378	37 (31.89)	48 (41.38)	0.02	0.875	20 (17.24)	65 (56.03)	0.903	0.342		
Female	11 (9.48)	20 (17.24)			14 (12.07)	17 (14.65)			10 (8.62)	21 (18.10)				
Tobacco consumption														
Yes	12 (10.34)	38 (32.76)	1.20	0.274	19 (16.37)	31 (26.72)	1.27	0.260	12 (10.34)	38 (32.76)	0.159	0.690		
No	22 (18.96)	44 (37.93)			32 (27.59)	34 (29.31)			18 (15.52)	48 (41.38)				
Alcohol consumption														
Yes	2 (1.72)	8 (6.89)	0.458	0.499	2 (1.72)	8 (6.89)	2.55	0.110	2 (1.72)	8 (6.89)	0.196	0.658		
No	32 (27.59)	74 (63.79)			49 (42.24)	57 (49.14)			28 (24.14)	78 (67.24)				
Grade														
Well	13 (11.21)	60 (51.72)	12.6	0.002*	28 (24.14)	45 (38.79)	2.53	0.283	12 (10.34)	61 (52.59)	9.19	0.010*		
Moderate	20 (17.24)	21 (18.10)			22 (18.96)	19 (16.38)			17 (14.65)	24 (20.69)				
Poor	1 (0.86)	1 (0.86)			1 (0.86)	1 (0.86)			1 (0.86)	1 (0.86)				
Primary tumour status														
T1	5 (4.31)	59 (50.86)	33.6	<0.001*	17 (14.65)	47 (40.52)	18.5	<0.001*	4 (3.45)	60 (51.72)	30.4	<0.001*		
T2	3 (2.59)	5 (4.31)			4 (3.45)	4 (3.45)			3 (2.59)	5 (4.31)				
T3	10 (8.62)	8 (6.90)			12 (10.34)	6 (5.17)			8 (6.89)	10 (8.62)				
T4	16 (13.79)	10 (8.62)			18 (15.52)	8 (6.89)			15 (12.93)	11 (9.48)				
Lymph node metastasis														
N0	11 (9.48)	55 (47.41)	12.9	0.005*	20 (17.24)	46 (39.65)	6.91	0.075	8 (6.89)	58 (50)	15.7	0.001*		
N1	14 (12.07)	13 (11.21)			16 (13.79)	11 (9.48)			13 (11.21)	14 (12.07)				
N2	8 (6.89)	13 (11.21)			14 (12.07)	7 (6.03)			8 (6.89)	13 (11.21)				
N3	1 (0.86)	1 (0.86)			1 (0.86)	1 (0.86)			1 (0.86)	1 (0.86)				
TNM stage group														
I+II	5 (4.31)	50 (43.10)	20.6	<0.001*	16 (13.79)	39 (33.62)	9.39	0.002*	4 (3.45)	51 (43.96)	18.9	<0.001*		
III+IV	29 (25)	32 (27.59)			35 (30.17)	26 (22.41)			26 (22.41)	35 (30.17)				

TABLE 2 (Continued)

Patients' characteristics	VM		χ^2	p value	Laminin-5 γ 2		χ^2	p value	VM and laminin-5 γ 2 dual expression		χ^2	p value
	Positive n (%)	Negative n (%)			Positive n (%)	Negative n (%)			Positive n (%)	Negative n (%)		
VM												
Positive	-	-	-	-	30 (25.86)	4 (3.45)	38.3	<0.001*	-	-	-	-
Negative	-	-	-	-	21 (18.10)	61 (52.59)			-	-	-	-

* Significant values i.e. $p < 0.05$ are denoted in bold.

period: 56 months, range: 16–60 months), 36 patients (31.03%) were dead due to local recurrence or metastasis after surgery and 62 (53.45%) patients were alive with the rest being lost to follow-up or died of other diseases unrelated to OSCC and they were considered as censored for further analysis. The Kaplan–Meier plot of DFS and OS in OSCC patients with differential status of VM, laminin-5 γ 2 and their dual existence have been shown in Figure 2. The follow-up data demonstrated that the mean DFS of VM-positive cohort and laminin-5 γ 2-positive cohort was significantly inferior to that of VM-negative (log rank = 92.052, $p < .001$) and laminin-5 γ 2-negative (log rank = 40.575, $p < .001$) cohort; VM–laminin-5 γ 2 double-positive cohort also had significantly worse DFS compared with the respective double-negative cohorts (log rank = 125.283, $p < .001$). The distinguishing DFS rate among these groups was also reflected with the OS time. The mean OS of VM-positive cohort, laminin-5 γ 2-positive cohort and VM–laminin-5 γ 2 double-positive cohort was significantly poorer than that of VM-negative (log rank = 80.363, $p < .001$), laminin-5 γ 2 negative (log rank = 45.209, $p < .001$) and VM–laminin-5 γ 2 double-negative (log rank = 114.464, $p < .001$) cohorts respectively. These findings interpreted that VM and laminin-5 γ 2 either individually or together are important indicators of DFS and OS in OSCC patients. Additionally, we have also found the significant difference in the Kaplan–Meier plot of DFS and OS in the OSCC patients with differential expression of MMP2 and of VM-MMP2 duality (Figure S3).

3.5 | Prognostic impact of paired VM–laminin-5 γ 2 positivity on disease-free and overall survival

Based on the significant findings of univariate analysis, indicating the significance of clinicopathological parameters such as tumour grade, primary tumour status, lymph node metastasis, TNM stage group and occurrence of VM independently, as well as in conjunction with the expression of laminin-5 γ 2 in DFS and OS, the multivariate Cox proportional hazard regression model was applied to assess their role as independent survival risk factors (Table 3). The multivariate analysis revealed that in addition to primary tumour status and lymph node metastasis, the occurrence of VM [hazard ratio (HR): 1.696; 95% CI :1.030–2.791; $p = .038$], positive expression of laminin-5 γ 2 (HR: 1.327; CI: 1.013–1.739; $p = .040$) and VM–laminin-5 γ 2 double-positive status (HR: 9.896; CI: 1.286–76.173; $p = .028$) were proved to be independent risk factors for DFS. Similar to DFS, the occurrence of VM (HR: 3.081; CI: 1.428–6.651; $p = .004$), positive expression of laminin-5 γ 2 (HR: 1.424;

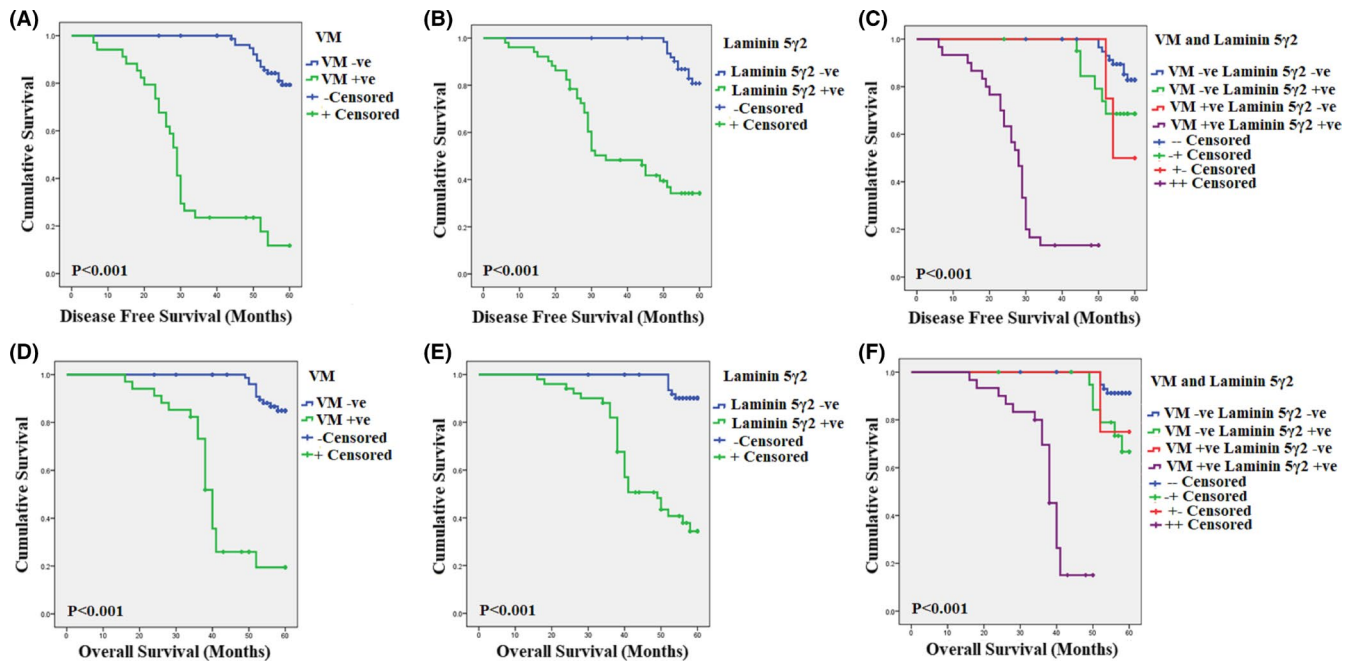


FIGURE 2 Kaplan–Meier analysis of the disease-free survival (DFS) and overall survival (OS) rate of patients with OSCC. DFS of patients in relation to (A) VM, (B) differential expressional status of laminin-5 γ 2, (C) VM and laminin-5 γ 2 dual status and OS of patients in relation to (D) VM, (E) differential expressional status of laminin-5 γ 2, (F) VM and laminin-5 γ 2 dual status

TABLE 3 Assessment of prognostic factors of disease-free survival (DFS) and overall survival (OS) by multivariate analysis of Cox proportional hazard model

Variables	Disease-free survival (DFS)				Overall survival (OS)			
	HR	<i>p</i> value	95% CI		HR	<i>p</i> value	95% CI	
			Lower	Upper			Lower	Upper
Grade	1.759	0.099	0.900	3.439	1.219	0.625	0.551	2.699
Primary tumour status	1.821	0.003*	1.221	2.717	1.559	0.054	0.992	2.449
Lymph node metastasis	2.944	<0.001*	1.632	5.311	2.865	0.002*	1.478	5.554
TNM stage group	0.307	0.123	0.068	1.379	0.494	0.486	0.068	3.589
VM only	1.696	0.038*	1.030	2.791	3.081	0.004*	1.428	6.651
Laminin-5 γ 2 only	1.327	0.040*	1.013	1.739	1.424	0.026*	1.043	1.945
VM and Laminin-5 γ 2 double positive	9.896	0.028*	1.286	76.173	21.401	0.033*	1.276	358.980
MMP2 only	1.091	0.582	0.801	1.485	1.084	0.697	0.721	1.630
VM and MMP2 double positive	1.047	0.951	0.238	4.619	0.282	0.232	0.035	2.251

* Significant values i.e. $p < 0.05$ are denoted in bold.

CI: 1.043–1.945; $p = .026$) and the simultaneous double-positive expression of VM–laminin-5 γ 2 (HR: 21.401; CI: 1.276–358.980; $p = .033$) were also found to be independent risk factor for OS. In support of the significant findings on the survival endpoints, the positive expression of MMP2 and its co-existence with VM were also considered for the analysis of risk factor assessment, but neither of them proved to be the independent prognostic factor for DFS and OS in our OSCC cohorts.

4 | DISCUSSION

Alternative vascularization influences the poor prognosis of cancer patients,^{53,54} evoking tumour resistance towards anti-angiogenic and anti-neoplastic therapy.⁵⁵ Vasculogenic mimicry is a leading pathological entity representing this state to which prompted us for a more comprehensive evaluation of VM and associated prognostic biomarkers underpinning in OSCC.



In this present study, we have evaluated the correlation of VM with expression of laminin-5 γ 2 in predicting the survival and prognosis of OSCC. We have inferred that the occurrence of VM is significantly prevalent in the poorly differentiated tumour with increased primary tumour size, higher lymph-node metastasis and TNM stage, which reflected the mechanistic link of VM to the invasion and metastasis attributing the aggressive and malignant progression of OSCC. Contextually, commonalities have been observed in other malignant tumours.^{56,57} Laminin-5 γ 2 complements ECM remodelling and is considered to be one of the most common downstream signalling proteins in molecular cascades associated with VM, that is TGF- β ,⁵⁸ VE-cadherin, EphA2, PI3-K⁵⁹ and MMP2.^{60,61} Moreover, laminin receptor integrin β 1-mediated FAK signalling has also been associated with VM-like network formation in human fibrosarcoma cells.⁶² On the contrary, the cooperative interaction between MMP2 and laminin-5 γ 2 has been well established in a number of malignancies including glioblastoma⁵⁸ and aggressive melanoma when cultured on a three-dimensional ECM.⁶³ Our study delineated a significant interrelation of VM, laminin-5 γ 2 and their coordinated alignment with the histological and conventional prognostic parameters such as tumour grade, primary tumour size, lymph node metastasis and TNM stage group highlighting the impact of integrating multiple facets of these markers that may benefit while assessing the risk factors in OSCC. The double positivity of vasculogenic mimicry–laminin-5 γ 2, as well as their individual positive expression, also had the significantly poorer DFS and OS in our study, which may act as a tool to predict a worse prognostic indication. Although a few recent studies indicated the individual prognostic significance of some VM-associated biomarkers including LGR5,⁶⁴ ALDH1, Beclin1, p16⁴⁹ and extracellular IL17-F,⁶⁵ the combinatorial approach of VM with its associated biomarkers^{14,66} is still a less explored area in OSCC. In this context, our investigation confirmed for the first time that both VM and laminin-5 γ 2 in combination provide better prognostic significance with higher statistical power including increased hazard ratio [(HR) = 9.896, p = .028 (DFS) and HR = 21.401, p = .033 (OS)] compared with individual expression of VM [(HR) = 1.696, p = .038 (DFS) and HR = 3.081, p = .004 (OS)] and laminin-5 γ 2 [HR) = 1.327, p = .040 (DFS) and HR = 1.424, p = .026 (OS)]. Collectively, these findings indicate the complementarity of VM and laminin-5 γ 2 as powerful risk factor for DFS and OS in OSCC. Our study also illustrated that in spite of being the upstream modulator of laminin-5 γ 2, MMP2 was not found to be performed as an independent risk factor in association with VM. We further validated the manual quantification data with the inputs from automatic profiler and showed a linear pattern.⁶⁷ Indeed,

digital quantitative pathology is an evolving modality and needs further validation before its routine adoption as stand-alone method. Knowing the therapeutic challenges of late refractory oral malignancies and roles of novel prognostic biomarkers in informed treatment decision, these findings will provide important contextual guidance for defining appropriate clinical strategies.

In conclusion, the study revealed that the expression of the extracellular matrix protein Laminin-5 γ 2 coordinated with VM is significantly associated with tumour grade, primary tumour size, lymph node metastasis and TNM stage. Co-expression of vasculogenic mimicry with laminin-5 γ 2 underlines the independent prognostic impact and correlates with the decreased disease-free and overall survival in OSCC patients. Further validation of these findings in large independent studies would provide important predictive opportunities for better guidance towards effective treatments.

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CONFLICT OF INTEREST

None.

AUTHOR CONTRIBUTIONS

DS and NM conceptualized and designed the study. DS and DM performed the experiments and analysed the data. NA and SS supplied tissue specimens. SMM validated the data and provided clinical insights. SM interpreted the data and performed statistical analysis. DS, BM and NM drafted the manuscript. BM and NM reviewed and edited the manuscript. All the authors approved the final version of the manuscript.

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