

PD-L1 Expression Confers Better Prognosis in Locally Advanced Oral Squamous Cell Carcinoma

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Abstract. *Background/Aim:* Clinical trials with therapies targeting immune checkpoint molecules have shown promising results in several tumor types. However, the predictive and prognostic values of these immunological factors for locally advanced oral squamous cell carcinomas (LAOSCC) remain unclear. Our purpose was to evaluate the expression and prognostic value of programmed cell death-ligand1 (PD-L1) and PD-L2 and to correlate their expression with the degree of infiltration by CD8⁺ cells in LAOSCC. *Patients and Methods:* A total of 84 patients with LAOSCC were included. PD-L1, PD-L2 and CD8 expression was detected in the tumor tissue using immunohistochemistry and was tested for correlation with clinical outcome. *Results:* PD-L1 and PD-L2 were expressed in 52.4% and 23.8% of LAOSCC cases, respectively. PD-L1 positivity was significantly associated with superior disease-free ($p=0.024$) and overall ($p=0.008$) survival of the patients and retained significance in multivariate analysis. PD-L1 positivity was correlated with CD8 density. *Conclusion:* PD-L1 expression was associated with CD8⁺ tumor-infiltrating lymphocytes and better outcome in patients with LAOSCC.

The recent demonstration that blockade of the programmed cell death-1/-ligand 1 (PD-1/PD-L1) or PD-L2 checkpoint pathway

is effective against several cancer types, including melanoma (1) and non-small cell lung cancer (2), has led to the clinical testing of inhibitors of PD-1 and its ligands in head and neck squamous cell cancers (HNSCC) (3). Immune check points such as PD-1 are manipulated by tumors to allow tumor growth that is unchecked by the immune system. Overexpression of PD-L1 by tumor cells activates the PD-1 and its ligands checkpoint pathway, by binding to the PD-1 receptor, and attenuating the immune response. The binding of antibodies targeting immune checkpoints such the binding of PD-1 to its ligands can ‘release the brakes’ and induce an immune response against the tumor. Solid cancers are infiltrated to varying degrees by immune cells [tumor-infiltrating lymphocytes (TILs)], and this phenomenon was suggested to be a manifestation of the host immune response against cancer cells (4). Given that PD-1 ligands on tumor cells interact with TILs in the tumor microenvironment, a comprehensive analysis of molecules related to PD-1 and its ligands might provide invaluable information for determination of the biological and clinical relevance of this pathway for head and neck cancer.

The present study aimed to evaluate the prognostic significance of PD-L1 and PD-L2 expression and their correlation with TILs in locally advanced oral squamous cell carcinoma (OSCC).

Patients and Methods

A total of 84 patients with primary advanced OSCC who underwent surgical resection at the Saitama Medical School International Medical Center (Saitama, Japan) from April 2007 to December 2014 were included in this study. Patients who had received induction chemotherapy or radiotherapy for the head and neck region before surgery or who exhibited distant metastasis at the time of diagnosis were excluded. This study was approved by the Institutional Review Board of Saitama Medical School International Medical Center (14-192).

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Key Words: PD-L1, PD-L2, immune checkpoint, prognosis, head and neck cancer.

Immunohistochemistry (IHC) of PD-L1, PD-L2 and CD8. IHC was performed to estimate the expression of PD-L1 and PD-L2, and the presence of CD8⁺ TILs in tumor samples. Tumor sections were deparaffinized with xylol and dehydrated in an alcohol series. The sections were washed in hydrogen peroxide, followed by distilled water washes. The slides were then subjected to heat-induced antigen retrieval using EDTA buffer (pH 9.0) at 100°C for 20 min, and were then washed in distilled water. Subsequently, the slides were incubated in a moist chamber, with the following specific primary antibodies: A rabbit monoclonal antibody to PD-L1 (clone SP142; Spring Bioscience, Pleasanton, CA, USA), a rabbit polyclonal antibody to PD-L2 (clone 80380; Sigma-Aldrich, Missouri, St. Louis, MO, USA), and a mouse monoclonal antibody to CD8 (clone C8/144B; ThermoFisher Scientific, Waltham, MA, USA), at 4°C overnight. The antibodies were diluted as follows: anti-PD-L1 and anti-CD8, 1:100; anti-PD-L2 1:10. The slides were then washed in phosphate-buffered saline (PBS), pH 7.4. The slides were subsequently incubated with primary anti-mouse or anti-rabbit antibody, followed by incubation with the labeled polymer, Envision™+ Dual Link System-HRP (DAKO, Santa Clara, CA, USA) using two sequential 30-min incubations. Staining was completed with a 5-10 min incubation with 3,3'-diaminobenzidine (DAB)+ substrate-chromogen (DAKO). Finally, the slides were washed in tap water, counterstained with hematoxylin for 30 s, dehydrated and mounted.

PD-L1 positivity of specimens was defined based on a 5% expression threshold, according to reported clinical studies (1, 5). The threshold of PD-L2 expression was defined as for PD-L1 for descriptive purposes. The number of TILs (CD8⁺ cells) was counted in at least four different high-power fields (hpf; ×40 objective, ×10 eyepiece) for each specimen. Fields with the most abundant TILs in a specimen were selected for counting. The average number of TILs of all specimens was used as the threshold level for determination of high or low TIL frequency for each specimen.

Statistical analysis. All statistical analyses were performed with EZR (Saitama Medical Center, Jichi Medical University, Saitama, Japan), which is a graphical user interface for R (The R Foundation for Statistical Computing, Vienna, Austria, ver. 2.13.0). More precisely, it is a modified version of R commander (ver. 1.6-3) designed to add statistical functions frequently used in biostatistics (6). Correlations between PD-L1 or PD-L2 expression and patient characteristics were analyzed using the chi-squared or Fisher's exact test for categorical variables. Overall survival (OS) and progression-free survival (PFS) were calculated by the Kaplan–Meier method; differences were assessed by a log-rank test. We performed univariate analyses using the Cox proportional hazards regression model to identify prognostic factors associated with OS and PFS, which were then considered in a multivariate Cox proportional hazards regression analysis. All *p*-values were two sided and *p*-values of 0.05 or less were considered statistically significant.

Results

Clinicopathological characteristics of the patients. Eighty-four patients (57 males (67.9%) and 27 females (32.1%)) were retrospectively included in the study. The median age of the patients at diagnosis was 68 years (range=20-92 years). The mean duration of follow-up was 40.6 months,

Table I. Clinicopathological characteristics of patients.

Characteristic	Value
Age, years	
Median	68
Range	20-92
Gender, n	
Male	57
Female	27
Follow-up period, months	
Median	40.6
Range	3.8-89.6
Smoking status, n	
Never-smoker	33
Smoker	39
Unknown	12
Performance status, n	
0	56
1	24
2	4
Clinical stage, n	
III	13
IVA	71
Subsite, n	
Cheek lining	5
Gingiva	31
Hard palate	4
Tongue	32
Floor of mouth	12

with a range of 3.8-89.6 months. Thirteen patients (15.5%) had stage III disease and 71 patients (84.5%) had stage IVA disease at the time of diagnosis. Tumor subsites and patient characteristics are shown in Table I.

The expression of PD-L1 and PD-L2 and infiltration of CD8⁺ TILs in locally advanced OSCC. Immunostaining of PD-L1 and PD-L2 was observed in the membrane or cytoplasm of tumor cells and stromal lymphocytes. Representative images of IHC staining of PD-L1, PD-L2 and CD8 are presented in Figure 1A-C, respectively. Expression of PD-L1 and PD-L2 was positive in 52.4% and 23.8% of tumors, respectively (Table II). PD-L1 expression was not significantly correlated with PD-L2 expression (Spearman's rank correlation; *r*=0.141, *p*=0.2). The number of TILs ranged from 0 to 353 per hpf (average 83/hpf TILs). We therefore divided the tumors into two groups based on the number of CD8⁺ TILs: a group with high TIL frequency, which had more than 83 TILs, and a group with low TIL frequency, with fewer than 83 TILs.

Comparative analysis of PD-L1 and PD-L2 expression and CD8⁺ TILs, and their association with clinicopathological characteristics in locally advanced

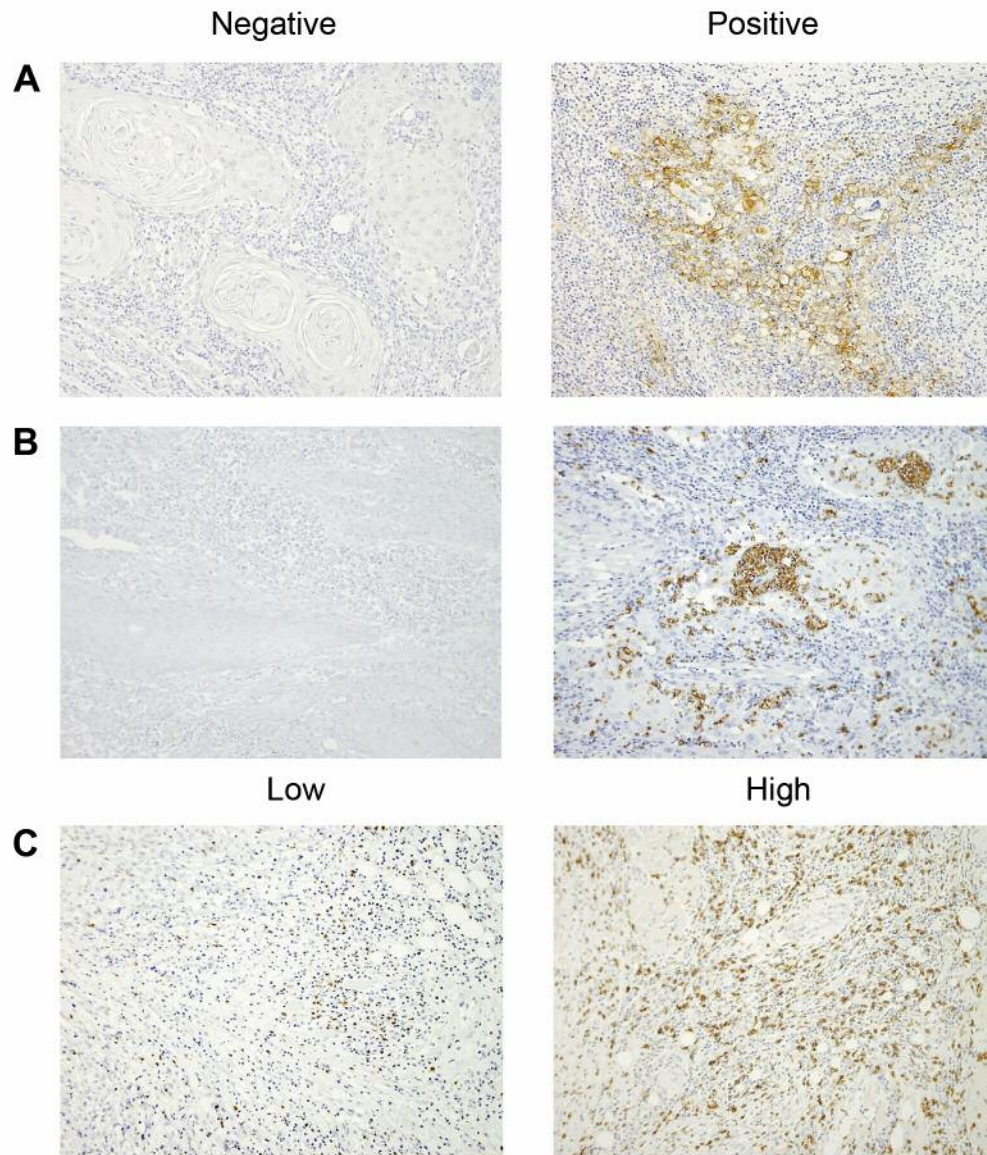


Figure 1. Representative expression patterns of programmed cell death-ligand1 (PD-L1), PD-L2, and CD8 in advanced oral squamous cell carcinomas (OSCC) samples. Representative negative and positive staining of samples of advanced oral SCC for PD-L1 (A) and PD-L2 (B), and CD8 staining of samples with low and high numbers of CD8⁺ tumor-infiltrating lymphocytes (TILs) (C) are shown. The number of CD8⁺ TILs was counted using at least four different high-power fields ($\times 40$ objective $\times 10$ eyepiece) that contained the highest level of TILs.

OSCC. The clinicopathological characteristics of patients with advanced OSCC, grouped according to their expression of PD-L1 and PD-L2, are shown in Table II. PD-L1 positivity was significantly positively correlated with the number of TILs ($p < 0.001$, Table II and Figure 2). In contrast, no significant relationship between PD-L2 expression and the number of TILs was detected. Moreover, significantly positive correlations between PD-L1 expression and recurrence ($p = 0.024$) and female sex

($p = 0.010$), and between PD-L2 and stage ($p = 0.011$) were detected (Table II).

PD-L1 expression represents an independent predictor of survival of patients with locally advanced OSCC. We next performed Kaplan–Meier and Cox proportional regression analyses to determine whether the TIL number or PD-L1/PD-L2 positivity correlated with PFS or OS of patients following treatment.

Table II. Clinicopathological characteristics of patients with oral squamous cell carcinoma, including cluster of differentiation 8-positive (CD8⁺) tumor-infiltrating lymphocytes (TILs), according to programmed cell death-ligand 1 (PD-L1) and PD-L2 expression in tumor.

	PD-L1 expression			PD-L2 expression		
	Negative	Positive	<i>p</i> -Value	Negative	Positive	<i>p</i> -Value
Total	40	44		64	20	
Gender, n						
Female	7	20	0.010	21	6	1.000
Male	33	24		43	14	
Age, years						
<65	15	13	0.492	22	6	0.792
≥65	25	31		42	14	
BMI, kg/m ²						
<18	7	6	0.764	10	3	1.000
≥18	32	38		53	17	
Smoking history, n ^a						
Never smoker	12	21	0.103	29	4	0.0508
Smoker	22	17		26	13	
Performance status, n						
0	25	31	0.492	45	11	0.277
1/2	15	13		19	9	
T Stage, n						
1/2	8	11	0.613	13	6	0.373
3/4	32	33		51	14	
N Stage, n						
0	11	13	0.734	20	4	0.449
1	6	9		10	5	
2	23	22		34	11	
Stage, n						
III	4	9	0.235	6	7	0.011
IV	36	35		58	13	
Recurrence, n						
No	22	33	0.024	42	13	0.866
L/R	6	8		10	4	
Distant	12	3		12	3	
PORT/POCRT, n						
Yes	20	16	0.271	26	16	0.605
No	20	28		38	10	
CD8 ⁺ TILs, n						
Low	29	13	<0.001	33	9	0.798
High	11	31		31	11	

BMI: body mass index; L/R: local/regional only; PORT/POCRT: postoperative radio therapy/chemoradiotherapy. ^aData missing for 13 patients.

Kaplan–Meier curves showed significantly poorer PFS and OS of patients with PD-L1-negative tumors compared to those with PD-L1-positive ones ($p=0.024$, and $p=0.008$, respectively). No significant differences in PFS and OS according to PD-L2 status were found. Such survival analysis also demonstrated that the number of TILs was not correlated with PFS or OS (log rank test: $p=0.35$, and $p=0.058$, respectively) (Figure 3). Univariate analysis (Cox regression model) showed that two variables, cN2 and PD-L1 positivity, were significantly correlated with PFS and OS (Table III). Multivariate analyses (Cox regression model) for PFS and OS of these factors showed that cN2 ($p=0.0324$ and $p=0.010$, respectively) and PD-L1

positivity ($p=0.0315$ and $p=0.008$, respectively) were significantly correlated with PFS and OS (Table III).

Finally, we performed sub-group analysis to examine the association of TIL frequency with PD-L1 expression, and its relationship with prognosis. We divided the tumors into four groups based on their positive or negative PD-L1 expression and on their high or low TIL frequency. The PD-L1-positive group with high TIL frequency displayed a tendency towards prolonged survival, whereas the PD-L1-negative group with high TIL frequency displayed a tendency towards shorter survival, although this tendency was not statistically significant ($p=0.672$, Figure 4).

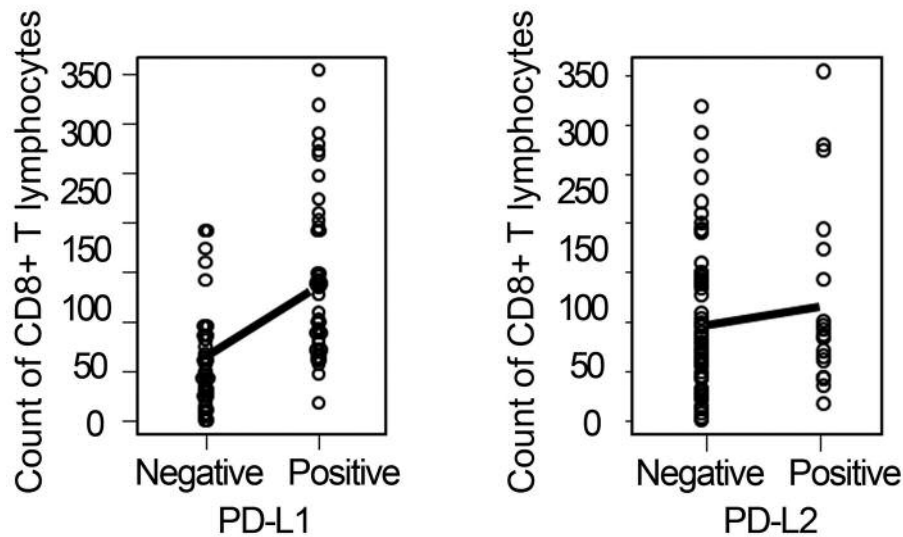


Figure 2. Correlation of programmed cell death-ligand1 (PD-L1) and PD-L2 expression and the number of CD8⁺ tumor-infiltrating lymphocytes (TILs). The CD8⁺ count was determined in PD-L1- and PD-L2-negative and -positive samples. The number of CD8⁺ TILs was significantly positively correlated with PD-L1 expression (A) (Spearman's $\rho=0.574$, $p<0.001$), but not with PD-L2 expression (B) (Spearman's $\rho=0.0818$, $p=0.459$).

Discussion

Although blockade of the PD-1 and PD-L1/PD-L2 pathway with monoclonal antibodies has recently emerged as a new therapeutic modality for head and neck cancer, the association between PD-L1 expression and prognosis in advanced HNSCC has remained largely controversial (7-9). In the present study, we examined PD-L1 and PD-L2 expression using IHC in patients with locally advanced OSCC who had undergone surgery, and found that patients who had positive expression of PD-L1 had significantly better survival than patients who were negative for PD-L1. Our results are not in agreement with previous studies that demonstrated an adverse prognostic effect of PD-L1 expression assessed by IHC in several cancer types including renal, colorectal and lung cancer (10-12). However, other studies in metastatic melanoma, non-small cell lung cancer, Merkel cell carcinoma, and laryngeal cancer, demonstrated a positive association of high PD-L1 expression and increased TIL frequency with longer survival (8, 13-15).

We also found a significant positive association between PD-L1 expression and the frequency of TILs. PD-L1 expression was reported to be up-regulated by CD8⁺ cell secretion of interferon gamma (IFN- γ) and patients with melanoma with TIL-rich tumors expressing PD-L1 are likely to have better immune surveillance (13). The prognostic importance of tumor infiltration with CD8⁺ cells has been demonstrated in breast, esophageal, lung, ovarian, colon, anal, and head and neck cancer (16, 17). Tumei *et al.* also demonstrated that the location of infiltrating CD8⁺ cells and PD-L1-expressing cells within a tumor specimen is of

particular importance; detection of CD8⁺ TILs and PD-L1-expressing cells at the invasive margin was associated with better response to therapy in melanoma (18). This finding suggests that CD8⁺ TILs have a crucial role in killing tumor cells when the PD-1/PD-L1 pathway is blocked. In our cases, the PD-1/PD-L1 pathway might have been blocked after surgery because almost all of the PD-L1 that was present in the local site was resected with the tumor. On the other hand, CD8⁺ TILs around the tumor were likely to remain at the local site after tumor resection. We speculate that the number of CD8⁺ TILs is increased in PD-L1-positive cases, and that the increase in CD8⁺ TILs was the reason why patients with PD-L1-positive tumors had a good prognosis.

Since PD-L1 positivity indicates the presence of higher numbers of TILs, anti-PD-1/PD-L1 therapy against such locally advanced OSCC would be effective *via* TILs. Consistent with this possibility, PD-L1 expression has been reported to be the most important predictor of the responsiveness of various cancer types to PD-L1 or PD-1 blocking antibodies (2, 19, 20).

It has already been shown that patients with HNSCC whose tumors contained infiltrating CD8⁺ TILs had better survival than those without TILs (8). Interestingly, in our study, PD-L1 positivity conferred a tendency for a more favorable influence on survival when high numbers of TILs were present in the specimen compared to when low numbers of TILs were present (Figure 4). These results together with our correlation study between PD-L1 positivity and numbers of TILs (Figure 2) suggest that PD-L1 may be a marker for the presence of TILs. In fact, TILs have been reported to stimulate PD-L1

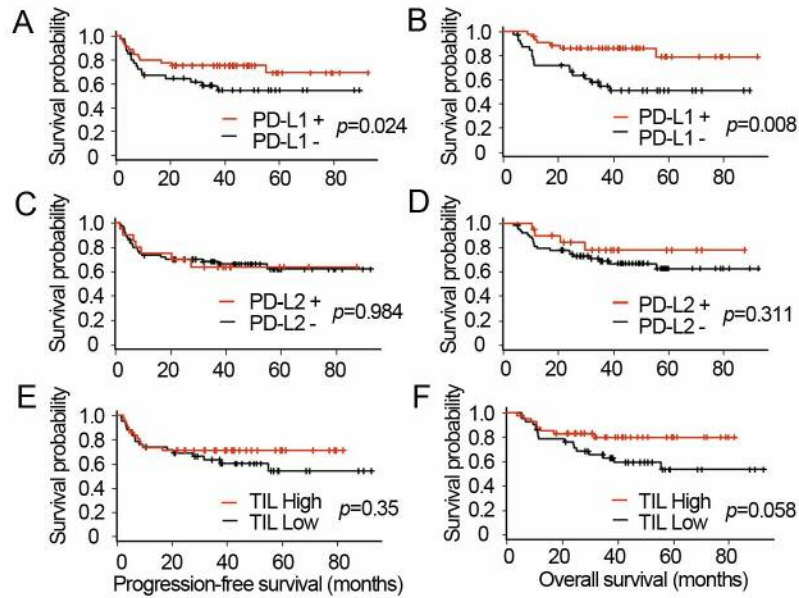


Figure 3. Correlation of programmed cell death-ligand1 (PD-L1) and PD-L2 expression and the number of CD8⁺ tumor-infiltrating lymphocytes (TILs) with survival. Kaplan–Meier curves for analysis of the effect of PD-L1 expression (A, B), PD-L2 expression (C, D) and the number of CD8⁺ CTLs (E, F) in tumor samples of patients with locally advanced oral squamous cell carcinomas on progression free survival (PFS) (A, C, E) and overall survival (OS) (B, D, F) are shown. *p*-Values were calculated using a log-rank test.

production by secreting IFN- γ in melanoma (13). In our study, there were 14 patients whose tumors were PD-L1-positive but which had low numbers of TILs (Figure 4). In such cases, PD-L1 expression is likely to be driven by pathways other than the IFN- γ pathway. The phosphoinositide 3-kinase/mitogen-activated protein kinase and zinc finger E-box-binding homeobox 1/miR-200 pathways have been reported to stimulate PD-L1 expression (21, 22). It is possible that the prognostic significance of PD-L1 may not be clear when PD-L1 expression is not driven by CD8⁺ cell-derived IFN- γ .

Our retrospective study had several limitations. Firstly, the sample size was relatively small. Secondly, PD-L1 and PD-L2 expression was evaluated using the antibodies of clone SP142 and clone 80380, respectively. There are many antibody clones for detection of PD-L1 and PD-L2, and it remains unclear which clones are the best for evaluation of the expression status of PD-L1 and PD-L2. Thirdly, positivity of PD-L1 and PD-L2 expression was determined based on a 5% expression threshold but this threshold is also controversial. Further studies are warranted to standardize the method of assessment of PD-L1 and PD-L2 expression by IHC in larger samples in head and neck cancer.

In conclusion, our study demonstrated that PD-L1 and PD-L2 were expressed in approximately half and one-quarter, respectively, of patients with locally advanced OSCC. Moreover, the expression of PD-L1 was positively associated with the frequency of TILs and better survival.

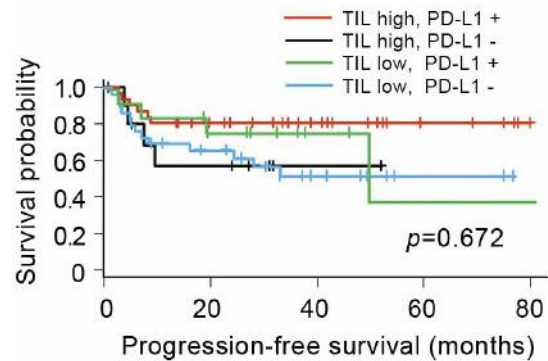


Figure 4. Kaplan–Meier curves of progression-free survival (PFS) of subgroups of patients with locally advanced oral squamous cell carcinomas grouped according to CD8⁺ count and programmed cell death-ligand1 (PD-L1) expression status. The *p*-value was calculated using a log-rank test. The CD8-high/PD-L1-positive group displayed a tendency towards better PFS than the other groups and the CD8-low, PD-L1-positive group displayed a tendency towards worse PFS than the other groups, although the differences between the groups were not significant.

Conflicts of Interest

The Authors declare that there is no conflict of interest in regard to this study.

Table III. Univariate and multivariate analyses of parameters associated with overall (OS) and progression free (PFS) survival.

Parameter	OS						PFS					
	Univariate analysis			Multivariate analysis			Univariate analysis			Multivariate analysis		
	HR	95% CI	p-Value	HR	95%CI	p-Value	HR	95% CI	p-Value	HR	95% CI	p-Value
≥65 years old	1.36	0.569-3.27	0.487				0.975	0.452-2.10	0.950			
Female gender	1.65	0.659-4.13	0.285				1.10	0.504-2.43	0.797			
BMI	0.521	0.193-1.40	0.197				0.706	0.268-1.85	0.480			
Smoker	0.857	0.543-1.36	0.511				0.691	0.307-1.55	0.373			
cT3/4	1.55	0.532-4.55	0.420				2.94	0.890-9.73	0.0766			
cN2	2.35	1.35-9.72	0.011	3.66	1.36-9.86	0.010	2.45	1.08-5.54	0.0313	2.439	1.078-5.520	0.0324
Margin-positive	2.09	0.622-7.07	0.233				1.36	0.411-4.52	0.610			
PD-L1-positive	0.257	0.102-0.649	0.006	0.256	0.101-0.646	0.008	0.576	0.274-0.956	0.0372	0.541	0.278-0.894	0.0315
PD-L2-positive	0.442	0.132-1.486	0.187				1.01	0.431-2.37	0.978			
CD8 high	0.499	0.214-1.16	0.109				0.717	0.342-1.50	0.378			

BMI: Body mass index; PD-L1/2: programmed cell death-ligand 1/2; CD8: cluster of differentiation 8; HR: hazard ratio; CI: confidence interval.

Acknowledgements

This study was supported by a grant from Saitama Medical University International Medical Center, H26 Hidaka Project (26-D-1-05) and was partly supported by a grant from Setsuro Fujii Memorial, The Osaka Foundation for Promotion of Fundamental Medical Research. The authors thank KW, NA and technicians in the Department of Pathology, Saitama Medical University International Medical Center, for their technical assistance.

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Received December 26, 2016

Revised February 7, 2017

Accepted February 8, 2017