


Prognostic impact of the tumor immune microenvironment in synovial sarcoma

Naoki Oike¹  | Hiroyuki Kawashima¹ | Akira Ogose^{1,2} | Tetsuo Hotta¹ | Hiroshi Hatano³ | Takashi Ariizumi¹ | Taro Sasaki³ | Tetsuro Yamagishi¹ | Hajime Umezu⁴ | Naoto Endo¹

¹Division of Orthopedic Surgery, Graduate School of Medical and Dental Sciences, Niigata University, Niigata, Japan

²Department of Orthopedic Surgery, Uonuma Kikan Hospital, Niigata, Japan

³Department of Orthopedic Surgery, Niigata Cancer Center Hospital, Niigata, Japan

⁴Department of Pathology, Graduate School of Medical and Dental Sciences, Niigata University, Niigata, Japan

Correspondence

Hiroyuki Kawashima, Division of Orthopedic Surgery, Graduate School of Medical and Dental Sciences, Niigata University, Niigata, Japan.

Email: inskawa@med.niigata-u.ac.jp

The association between the immune status within the tumor microenvironment and prognosis in synovial sarcoma is not well understood. We aimed to investigate the tumor immune microenvironment and analyze its prognostic impact for patients with synovial sarcoma. A total of 36 primary patients who were treated in our institution were retrospectively evaluated. Infiltration of lymphocytes (CD4+, CD8+, and FOXP3+), CD163+ macrophages, and expression of human leukocyte antigen (HLA) class I and programmed death ligand 1 (PD-L1) were evaluated by immunohistochemistry. Moreover, we investigated PD-L1 and programmed death ligand 2 (PD-L2) mRNA expression in 19 of the 36 cases, using real-time PCR. The Kaplan-Meier method was used to estimate overall survival and progression-free survival. Infiltration of lymphocytes and macrophages varied among the patients. Furthermore, the expression of HLA class I was negative or downregulated in 11 specimens. No PD-L1 expression was observed using immunohistochemistry. Moreover, although PD-L1 mRNA expression was observed in 18 of 19 specimens, the expression level was low. A higher infiltration of CD8+ or FOXP3+ lymphocytes in patients was associated with a favorable overall survival. In addition, a higher infiltration of CD163+ macrophages indicated a significantly worse overall and progression-free survival. Infiltration of CD4+ lymphocytes, HLA class I, PD-L1, and PD-L2 expression were not associated with patient prognosis. This represents the first report investigating the tumor immune microenvironment as a prognostic factor in synovial sarcoma, indicating that CD163+ macrophages are associated with tumor progression. Our results underscore the clinical significance of the tumor immune microenvironment in synovial sarcoma.

KEYWORDS

CD163+ macrophage, prognosis, programmed death ligand 1, synovial sarcoma, tumor immune microenvironment

Abbreviations: HLA, human leukocyte antigen; NY-ESO-1, New York esophageal squamous cell carcinoma 1; OS, overall survival; PD-1, programmed cell death 1; PD-L1, programmed death ligand 1; PD-L2, programmed death ligand 2; PFS, progression-free survival; SS, synovial sarcoma; TILs, tumor infiltrating lymphocytes; Treg, regulatory T cell.

1 | INTRODUCTION

Synovial sarcoma is a mesenchymal tumor that comprises 5%-10% of soft tissue sarcomas. The 5-year OS rate of synovial sarcoma is 64%-

This is an open access article under the terms of the Creative Commons Attribution-NonCommercial-NoDerivs License, which permits use and distribution in any medium, provided the original work is properly cited, the use is non-commercial and no modifications or adaptations are made.

© 2018 The Authors. *Cancer Science* published by John Wiley & Sons Australia, Ltd on behalf of Japanese Cancer Association.

76%.^{1,2} Standard treatment for primary localized SS consists of surgery, radiotherapy, and chemotherapy. However, local or metastatic recurrences occur in approximately 50% of patients at 5 years,³ and the benefit of adjuvant chemotherapy in localized primary SS cases is still controversial.² In addition, the prognosis of metastatic or relapsed SS remains poor. Thus, to improve the survival rate of patients with SS, the development of novel therapies is needed.

Synovial sarcoma presents a specific chromosomal translocation t(X:18)(p11.2;q11.2), resulting in SS18-SSX1 or SS18-SSX2, and rarely into the SS18-SSX4 fusion proteins.^{4,5} Moreover, the NY-ESO-1 cancer testis antigen is expressed in approximately 80% of SS cases.^{6,7} Recently, immunotherapies consisting of SYT-SSX or NY-ESO-1-delivered peptide vaccines^{8,9} or autologous T cells transduced with a T-cell receptor directed against NY-ESO-1 were attempted for patients with SS.¹⁰ Importantly, clinical trials using an adoptive cell transfer of autologous T cells transduced with anti-NY-ESO-1 have shown promise in triggering objective responses in patients with metastatic or refractory disease.¹⁰ These data suggest that immunotherapies may be effective in the treatment of SS.

Programmed cell death 1 and its ligands, PD-L1 and PD-L2, play a key role in the immune escape of tumor cells by inactivating T-cell function. Several authors have shown that antibodies blocking PD-1 or PD-L1 lead to improved clinical outcomes in a variety of malignancies.¹¹⁻¹³ However, previous reports have indicated that expression of PD-L1/PD-L2 has different prognostic value in various cancers.¹⁴⁻¹⁷ Association between PD-L1 expression and the prognosis of soft tissue sarcoma, including SS, was also examined. However, because these studies comprised a small number of patients with SS, the expression of PD-L1 in SS and its clinical significance remain controversial.¹⁷⁻²¹

Understanding how the tumor immune microenvironment relates to the prognosis of patients with SS may clarify the potential for effective immunotherapy for the treatment of SS. However, to the best of our knowledge, the tumor immune microenvironment including TILs, macrophages, and the expression of HLA class I and PD-L1 are not well understood in SS. The purpose of the present study was to investigate the tumor immune microenvironment and evaluate the prognostic impact of various immunological factors in patients with SS. In addition, we are the first to investigate the tumor immune environment of primary non-treated patients with SS and analyze its clinical significance in SS.

2 | MATERIALS AND METHODS

2.1 | Patient tissue samples

A total of 39 primary patients with SS, who were diagnosed and treated in Niigata University Hospital and Niigata Cancer Center Hospital between 1983 and 2017, were enrolled. To evaluate the natural tumor immune microenvironment, three patients were excluded because they had already received some treatment in other hospitals or for which specimens without neoadjuvant chemotherapy were not available. Thus, a total of 36 patients were retrospectively examined in this study. We used biopsy specimens for the immunohistochemical examination if the patient had

received neoadjuvant chemotherapy. In any other case, we used surgical specimens for the examination. In all cases, H&E staining was prepared from the archives. The SS diagnosis was confirmed and representative samples, including viable tumor cells, were selected by an experienced pathologist. The patient's clinical information and follow-up data were collected from the medical records for statistical analysis. Clinical characteristics of the patients are summarized in Table 1. There were 23 male and 13 female patients with a mean age of 36 years (ranging from 10 to 76 years). Three of these patients showed lung metastasis at diagnosis. Ten tumors were located in the upper extremities, 23 in the lower extremities, and three in the trunk. The American Joint Committee on Cancer (AJCC) 7th edition staging system was used to classify disease staging;²² 14 patients were stage II, 19 were stage III, and three were stage IV.

This study was approved by the Institutional Review Board of the Niigata University Hospital (No. 2016-0024) and was conducted in line with the Declaration of Helsinki. Written informed consent was obtained from all patients and/or their families prior to their participation in this study.

2.2 | Immunohistochemistry

To evaluate the tumor immune microenvironment in patients with SS, immunohistochemical staining for TILs (CD4, CD8, FOXP3),

TABLE 1 Patient demographics and clinicopathological characteristics

Characteristic	n (%)
Total number of patients	36 (100)
Age at diagnosis, y	
<20	6 (16.7)
≥20	30 (83.3)
Gender	
Male	23 (63.9)
Female	13 (36.1)
Location	
Upper extremities	10 (27.8)
Lower extremities	23 (63.9)
Trunk	3 (8.3)
Tumor size, cm	
>5	22 (61.1)
≤5	14 (38.9)
Translocation	
SYT-SSX 1	24 (66.7)
SYT-SSX 2	6 (16.7)
NA	6 (16.7)
Stage (AJCC system)	
II	14 (38.9)
III	19 (52.8)
IV	3 (8.3)

AJCC, American Joint Committee on Cancer systems 7th edition staging manual; NA, not available.

CD163+ macrophages, HLA class I, and PD-L1 was conducted. For each specimen, the immunohistochemical analysis was carried out on 4- μ m sections from representative blocks of formalin-fixed, paraffin-embedded tumor tissues. The sections were deparaffinized with xylene and rehydrated with graduated ethanol. Heated antigen retrieval was carried out in citrate pH 6.0 (for CD163, HLA class I) or in Tris-EDTA pH 9.0 (Nichirei Bioscience, Tokyo, Japan, for CD4, CD8, FOXP3), using an autoclave (at 121°C) for 5 minutes. For PD-L1, the sections were boiled for 40 minutes in Tris-EDTA pH 9.0 (Nichirei Bioscience) for antigen retrieval. Endogenous peroxidase activity in tissues was blocked by incubation in 0.3% hydrogen peroxidase, and the sections were incubated with primary antibodies. The following monoclonal antibodies were used: CD4 (1F6, 1:50; Nichirei Bioscience), CD8 (C8/144B, 1:200; Nichirei Bioscience), FOXP3 (236A/E7, 1:200; Abcam, Cambridge, UK), CD163 (10D6, 1:500; Leica, Wetzlar, Germany), HLA class I (EMR8-5, 1:500; Hokudo, Sapporo, Japan), PD-L1 (28-8, 1:200; Abcam). The sections were incubated with the primary antibodies at 4°C overnight (CD4, CD8, CD163, FOXP3, and HLA class I), or at room temperature for 1 hour (PD-L1), followed by incubation with Histofine Simple Stain MAX PO MULTI (Nichirei Bioscience) at room temperature for 30 minutes. Peroxidase activity was detected with 3'-diaminobenzidine tetrahydrochloride (Simple Stain DAB; Nichirei Bioscience). Sections were counterstained with hematoxylin (Vector Laboratories Inc., Burlingame, CA, USA). Appropriate positive and negative controls were used for CD4, CD8, FOXP3, CD163, and PD-L1. Expression of HLA class I within endothelial cells served as an internal positive control. To validate our immunohistochemistry technique for PD-L1, immunohistochemical staining for PD-L1 MDA-MB231 was conducted.

2.3 | Evaluation of immunohistochemistry

For TILs and macrophages, tumor sections were evaluated at low magnification, and the areas with the most abundant TILs or macrophages were selected. A maximum of five high-power fields ($\times 200$) was then digitally photographed in a size of 0.093 mm² with an Olympus DP73 digital camera (Olympus, Tokyo, Japan) and counted manually. The count was conducted two times by an experienced pathologist who was blinded to previously available results and clinical information. Average value in an area of 1 mm² was used for the statistical analysis. For each immunohistochemical marker, we divided the patients into subgroups. The patients were classified into “high” and “low” infiltration groups except for HLA class I, where the “high” group displayed a number of TILs or macrophages that was above the median. For HLA class I, we classified the 36 cases into high (number of positive cells >50%), low (number of positive cells <50%), and negative (the samples were negative for these cells). For PD-L1, the patients were assigned to the “high” group when the number of positive cells >1%, as previously reported.^{21,23}

2.4 | Cell culture

HS-SYII and SYO-1 SS cells lines were obtained from Dr H. Sonobe (Department of Pathology, Kochi Medical School, Kochi, Japan)²⁴

and Dr A. Kawai (Department of Orthopedic Surgery, Okayama University, Okayama, Japan),²⁵ respectively. MDA-MB-231 breast cancer cell line was purchased from the European Collection of Authenticated Cell Cultures (ECACC, Salisbury, UK). All cell lines were cultured in DMEM (Invitrogen Life Technologies, Carlsbad, CA, USA) supplemented with 10% FBS and 1% Antibiotic-Antimycotic (Invitrogen Life Technologies). Cell lines were then incubated in a humidified atmosphere containing 5% CO₂ at 37°C.

2.5 | Quantitative real-time PCR

Quantitative real-time PCR for PD-L1 and PD-L2 was carried out in two SS cell lines and freshly frozen specimens from 19 non-treated patients with SS. Total RNA was isolated from cell lines or frozen specimens using ISOGEN (Nippon Gene, Tokyo, Japan) according to the manufacturer's protocol. Total RNA was converted to cyclic DNA using the PrimeScript RT reagent kit (TaKaRa Bio, Tokyo, Japan) according to the manufacturer's protocol. Quantitative real-time PCR was carried out using the SYBR Premix EX Taq II (Tli RNaseH Plus; TaKaRa, Shiga, Japan), and the results were analyzed using the Thermal Cycler Dice Real Time System TP800 (TaKaRa, Shiga, Japan). The *GAPDH* housekeeping gene was used as a reference gene. The primer sequences used were as follows: primer pairs used for human PD-L1 5'-CAATGTGACCAGCACACTGAGAA-3', (forward) and 5'-GGCATAATAAGATGGCTCCCAGAA-3' (reverse); PD-L2 5'-ATCCAACCTGGCTGCTTCCAC-3', (forward) and 5'-CTCCCAAGACCA CAGTTCA-3' (reverse); and *GAPDH*, 5'-GCACCGTCAAGGCTGA GAAC-3' (forward) and 5'-TGGTGAAGACGCCAGTGA-3' (reverse). mRNA expression levels in each sample were normalized by the *GAPDH* levels. Gene copy number for PD-L1 was calculated using a standard curve that was constructed using the MDA-MB-231 cell line. MDA-MB-231 was previously reported as a positive control and was used as a calibrator.²³ Nineteen patients were divided into two subgroups for the statistical analysis. The patients were considered as belonging to the “high” group when the PD-L1 or PD-L2 mRNA level was above the median, and the levels for the rest of the patients were low.

2.6 | Western blot analysis

HS-SYII and SYO-1 were subjected to western blot analysis to assess endogenous expression of the PD-L1 protein as previously reported.²⁶ Briefly, the cells were suspended in a sample buffer containing 62.5 mmol/L Tris, pH 6.8; 2% SDS; 5% glycerol; and 6 mol/L urea. Then, dithiothreitol (1 mol/L; Sigma-Aldrich, St Louis, MO, USA) and bromophenol blue were added to the lysates, and the lysates were boiled for 5 minutes and stored at -80°C. Equal amounts of proteins (50 μ g/lane) were run on 10% SDS-PAGE gels (Bio-Rad, Hercules, CA, USA) and were transferred to nitrocellulose membranes (GE Healthcare, Little Chalfont, Buckinghamshire, UK). The membranes were incubated overnight at 4°C with primary antibody against PD-L1(28-8, 1:2000; Abcam) and actin (AC-74, 1:3000; Sigma-Aldrich), and with HRP-labeled anti-rabbit IgG (GE Healthcare)

at room temperature. Bands were visualized by ECL detection reagents (GE Healthcare).

2.7 | Statistical analysis

Statistical analyses were conducted using IBM SPSS version 21.0 (IBM, New York, NY, USA) or GraphPad Prism v6.0 software (La Jolla, CA, USA). Spearman's rank correlation coefficient test was used to analyze the relationship between groups. OS and PFS were measured from the date of the initial biopsy. A terminal point for OS was determined as the time of death or the time the patient was last seen. A terminal point for PFS was defined as the time of local recurrence, appearance of distant metastasis, disease progression, or the last follow up. Kaplan-Meier method was used to estimate OS and PFS probabilities and survival differences were analyzed using the log-rank test. Multivariate analysis was conducted using the COX proportional hazard model. *P*-value < 0.05 was considered statistically significant.

3 | RESULTS

3.1 | Clinical course of the patients

Mean follow-up times were 112.7 months (6-366 months). OS probabilities of the 36 patients with SS at 5 and 10 years were 82.7% and 75.2%, respectively. The 5 and 10-year PFS probabilities were both 67.5%. Local recurrence was observed in one case and 12 patients developed distant metastasis. Sites of distant metastasis were the lung in 10 patients, and the lymph nodes, kidney, and vertebra in one patient.

3.2 | Infiltration of lymphocytes and macrophages

Representative images displaying high or low levels of infiltration for CD8+, FOXP3+ lymphocytes, or CD163+ macrophages are shown in Figure 1. Figure 2 shows the number of CD4+, CD8+, and FOXP3+ TILs, and CD163+ macrophages that infiltrated into the tumor per mm². Although CD163+ macrophages were seen in all patients, 10 patients were negative for CD4, one patient was negative for CD8, and nine patients were negative for FOXP3. Median number of infiltrating TILs and CD163+ macrophages was as follows: CD4+ lymphocytes 45.2 (0-697.3), CD8+ lymphocytes 188.3 (0-1358.0), FOXP3+ lymphocytes 18.3 (0-639.2), and CD163+ macrophages 444.4 (139.9-1628.1). Next, the correlation between infiltration of each immune cell was examined, and there was a correlation between the density of CD8+ and FOXP3+ lymphocytes and that of CD163+ macrophages. In addition, there was a strong correlation between CD8+ and FOXP3+ lymphocytes (Table 2).

3.3 | Expression of HLA class I and PD-L1

Immunohistochemical staining for HLA class I was conducted and representative images for the high or low expression of HLA class I are shown in Figure 1. Twenty-five (69.4%) of the 36 specimens

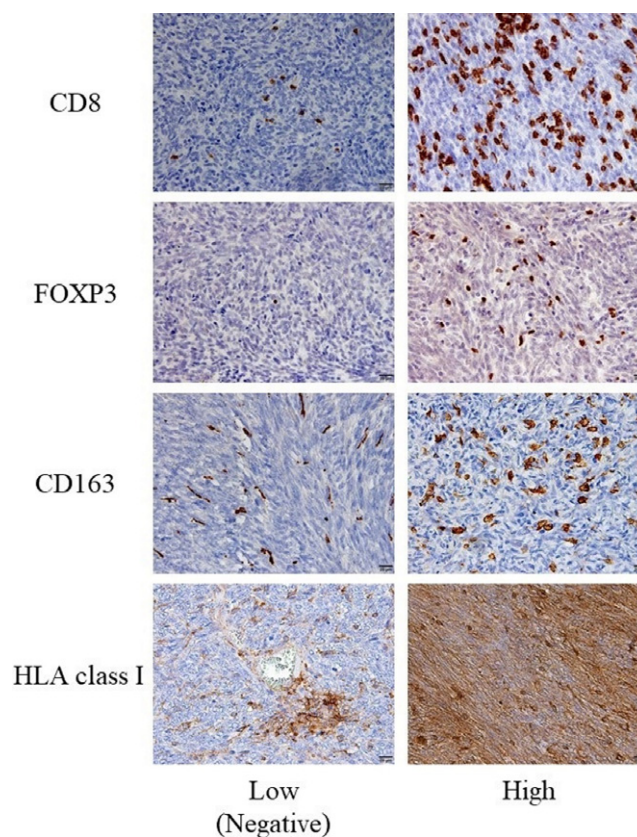


FIGURE 1 Representative images of immunohistochemical staining for CD8+ tumor-infiltrating lymphocytes (TILs), FOXP3+ TILs, CD163+ macrophages, and human leukocyte antigen (HLA) class I in synovial sarcoma (200 \times). (Left) Representative images of CD8+ TILs, FOXP3+ TILs, CD163+ macrophages, and HLA class I classified as “low” expression. (Right) Representative images of CD8+ TILs, FOXP3+ TILs, CD163+ macrophages, and HLA class I classified as “high” expression. Scale bar represents 20 μ m

were graded as high expression, five (13.9%) were classified as low expression, and the remaining six cases (16.7%) were considered negative for HLA class I. Patients with high expression of HLA class I showed a trend toward a higher number of infiltrated CD8+ lymphocytes compared with low or negative expression of HLA class I, but this was not statistically significant (data not shown). Although no case was positive for PD-L1 following the immunohistochemical analysis, intense staining for PD-L1 in MDA-MB231 cells was observed (Figure 3A).

3.4 | Programmed death ligand 1/PD-L2 mRNA and protein expression

Programmed death ligand 1 and PD-L2 mRNA expression were quantified in patients with SS and two SS cell lines by real-time PCR. Both synovial sarcoma cell lines and all 19 specimens expressed PD-L1 mRNA to a certain extent. However, no cases or cell lines showed a higher expression of PD-L1 mRNA than MDA-MB-231 (Figure 3B). Western blot analysis showed no protein expression of PD-L1 in both synovial sarcoma cell lines (Figure 3C). Therefore, PD-

FIGURE 2 Number of infiltrating CD4+, CD8+, FOXP3+ lymphocytes and CD163+ macrophages in the tumor. Graph shows number of lymphocytes and CD163+ macrophages per square millimeter ($\times 200$). CD163+ macrophages were observed in all patients. Error bars indicate the standard deviations

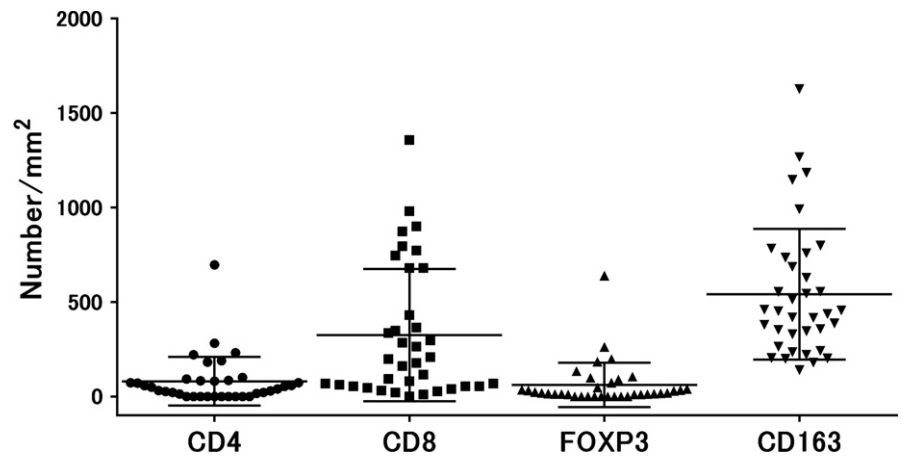


TABLE 2 Correlation of each immune parameter

	CD4 density	CD8 density	FOXP3 density	CD163 density
CD4 density		$r = 0.343$ $P = 0.040$	$r = 0.457$ $P = 0.005$	$r = 0.110$ $P = 0.525$
CD8 density			$r = 0.883$ $P < 0.001$	$r = 0.393$ $P = 0.018$
FOXP3 density				$r = 0.401$ $P = 0.015$
CD163 density				

r , correlation coefficient. Bold values are statistically significant ($P < 0.05$).

L1 mRNA expression of both cell lines was considered negative. Regarding PD-L2, five of 19 specimens showed a higher expression of PD-L2 mRNA than MDA-MB231 (Figure 3D). The expression level of both PD-L1 and PD-L2 was not correlated with the density of each immune cell and correlation between number of infiltrated CD163+ macrophages and expression level of PD-L1 and PD-L2 is shown in Figure 4.

3.5 | Prognostic significance of the immune microenvironment

Patient survival curves are shown in Figure 5. Patients with higher levels of CD8+ or FOXP3+ lymphocyte infiltration or lower levels of CD163+ macrophage infiltration were associated with a favorable OS. High expression of HLA class I showed a tendency toward longer OS compared with negative expression of HLA class I, but did not achieve statistical significance ($P = 0.052$). Next, to examine whether the tumor immune microenvironment was associated with the progression of SS, we evaluated PFS. Although no significant association was found for CD8+ lymphocytes or FOXP3+ lymphocytes, lower CD163+ macrophage infiltration levels in patients were also significantly associated with a longer PFS. These results suggest that CD163+ macrophages are associated with tumor progression in patients with SS. No prognostic

impact on OS and PFS was found for CD4+ lymphocyte infiltration, or HLA class I PD-L1, and PD-L2 expression. Univariate analysis of clinicopathological factors showed that advanced AJCC stage (III vs IV) was significantly associated with unfavorable prognosis (Table 3). Multivariate analysis was also carried out to evaluate the tumor immune microenvironment and clinicopathological factors. We excluded tumor size because AJCC staging system includes tumor size. We also excluded infiltration of FOXP3+ lymphocytes, because there was a strong correlation between the number of CD8+ and FOXP3+ lymphocytes. Multivariate analysis indicated that AJCC stage, number of CD8+ lymphocytes, and number of CD163+ macrophages were significantly associated with OS, respectively (Table 4).

4 | DISCUSSION

Some studies have examined the tumor microenvironment, including infiltration of CD8+ lymphocytes and expression of PD-L1 in soft tissue sarcomas.^{18,21,27} However, those studies often lump together heterogeneous types of soft tissue sarcomas and include limited numbers of patients with SS. To the best of our knowledge, the present study is the first report to focus on the tumor immune environment in patients with SS. We investigated the tumor immune microenvironment including infiltration of CD4+, CD8+, and FOXP3+ TILs, that of CD163+ macrophages, expression of HLA class I and PD-L1, and their association with the prognosis of SS.

Since CD8+ lymphocytes can kill tumor cells presenting tumor-associated antigen, CD8+ lymphocytes are essential for adaptive tumor immunity. As a result, a high infiltration of CD8+ TILs has a positive impact on the clinical course of many types of cancer.²⁸ However, studies investigating the associations between TILs and the prognosis of soft tissue sarcomas, including SS, are limited. Fujii et al²⁹ reported that a higher infiltration of CD8+ lymphocytes was associated with a favorable prognosis and correlated with distant metastasis-free survival in patients with angiosarcoma. In contrast, van Erp et al¹⁸ suggested that a high infiltration of CD8+ lymphocytes in SS tumors was associated with a worse metastasis-free

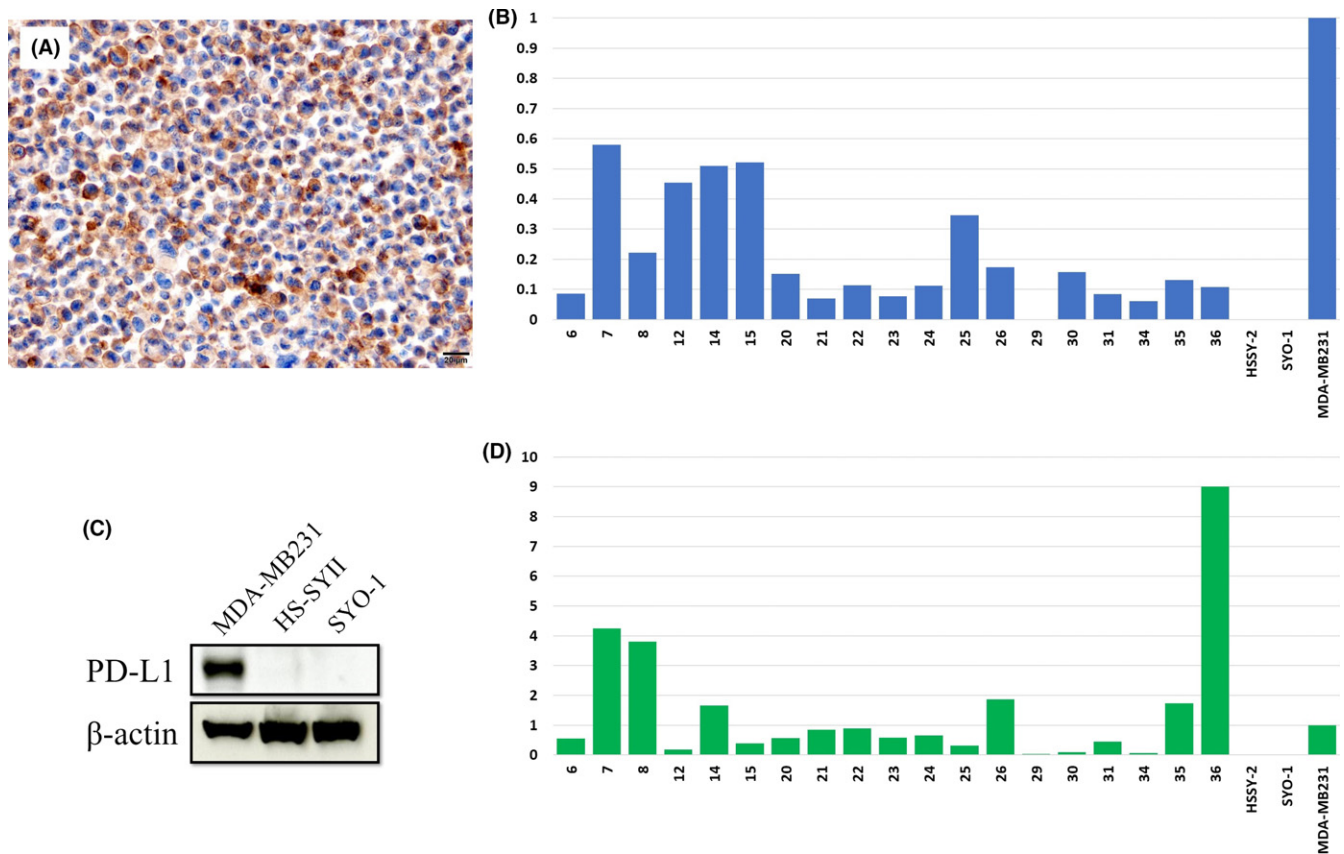


FIGURE 3 Expression of programmed death ligand 1 (PD-L1) and programmed death ligand 2 (PD-L2) in synovial sarcoma cell lines and tumor specimens. A, Immunohistochemical method for PD-L1 was validated using MDA-MB-231 breast cancer cell line. Intense staining for PD-L1 cells is observed. B, PD-L1 expressing MDA-MB-231 was used for calibration. Results for each specimen were standardized to the expression of *GAPDH*. mRNA PD-L1 expression was highest in MDA-MB-231. C, Western blot analysis of the expression of PD-L1 in synovial sarcoma cell lines show that PD-L1 protein was not detected in either HSSY-II or SYO-1. MDA-MB-231 was used as a positive control. D, PD-L2 expressing MDA-MB-231 was used for calibration. Five of 19 specimens showed higher mRNA PD-L2 expression than MDA-MB-231

survival in patients with SS. However, the OS of patients with SS was not clarified in this study, and the prognostic value of CD8 for patients with SS remains unclear. In addition, Nowicki et al¹⁹ reported that the density of intratumor CD8+ lymphocytes was not associated with OS or PFS in patients with SS. These results conflict with the data in our study. Our results indicate that although the infiltration of CD8+ lymphocytes is not associated with PFS, a higher infiltration of CD8+ lymphocytes is significantly associated with a favorable OS. Taken together, these findings suggest that although patients with a higher infiltration of CD8+ lymphocytes may develop distant metastases, the prognosis of these patients may not be poor.

Downregulation of HLA class I was associated with poor CD8+ infiltration and unfavorable prognosis.³⁰ However, the present study has shown that HLA class I expression is not associated with prognosis. This discrepancy might arise from the small number of samples in this study because negative expression of HLA class I showed a trend toward lower CD8+ infiltration and unfavorable prognosis.

Regulatory T cells represent a subset of CD4+ lymphocytes characterized by coexpression of CD4 and CD25.³¹ FOXP3, a nuclear transcription factor, functions as a master regulator for the

development of Treg and is considered the most reliable Treg marker.³¹ FOXP3+ Treg promote the evasion of cancer cells from immune responses.^{31,32} Although CD4+ lymphocytes did not show any association with the prognosis of patients with SS, in our study, a higher infiltration of FOXP3+ lymphocytes was significantly associated with a better OS. However, Que et al³³ evaluated the infiltration of FOXP3+ lymphocytes in 163 cases of soft tissue sarcoma, including 21 cases of SS, and showed that a higher infiltration of FOXP3+ lymphocytes was associated with a poor prognosis. This result is markedly different from ours. One possible explanation may be that as FOXP3+ T cells comprise functionally different subsets, including non-Treg, FOXP3 is not necessarily a specific marker of regulatory T cells.³⁴ Considering that Que's study did not distinguish SS from other soft tissue sarcomas, FOXP3+ lymphocytes that infiltrate into the tumor microenvironment in SS might represent a subset other than Treg. However, our data should be interpreted with care. As nine patients were negative for FOXP3, and the median value of infiltrated FOXP3+ lymphocytes was very low, a favorable prognosis might arise as a result of our choice of cutoff value. Although present study used a median value as the cutoff, Que et al defined "high infiltration" as infiltration occurring in greater than 5%

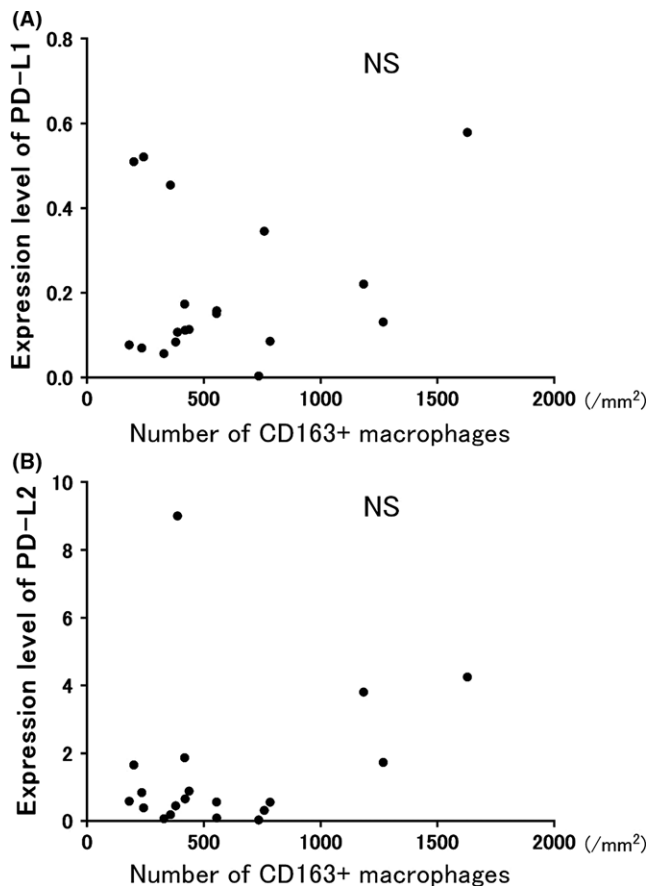


FIGURE 4 Correlation between CD163+ macrophages and programmed death ligand 1 (PD-L1) and programmed death ligand 2 (PD-L2) mRNA expression. There was no correlation between the number of infiltrated CD163+ macrophages and expression level of (A) PD-L1 and (B) PD-L2. NS, not significant

of tumor stromal cells. This discrepancy might influence the results of our study. Thus, further studies are needed to define appropriate thresholds.

Although some authors reported that PD-L1 expression was associated with the clinical prognosis of soft tissue sarcomas, including SS cases, expression of PD-L1 and the prognostic significance of PD-L1 in SS is still controversial.^{18-21,35} Kim C et al evaluated PD-L1 expression in 82 soft tissue sarcomas, including 19 SS, and Kim JR et al investigated 105 cases of soft tissue sarcoma, including 16 SS, by immunohistochemistry.^{20,35} They reported that 10 out of 19 (53%) and 12 out of 16 (75%) patients with SS were positive for PD-L1. In addition, Nowicki et al¹⁹ examined 29 SS specimens and showed that PD-L1 was expressed in SS. Conversely, D'Angelo et al²¹ investigated the expression of PD-L1 in 50 cases of soft tissue sarcoma, including five cases of patients with SS, whereas van Erp et al¹⁸ examined PD-L1 expression in a large cohort of primary sarcoma ($n = 202$), including 22 cases of patients with SS. Neither study identified any case that was positive for PD-L1 by immunohistochemistry. This discrepancy may possibly be as a result of the use of different antibodies or the definition of the cutoff value. Moreover, in the Nowicki et al study, greater than half of the specimens were metastatic specimens, and PD-L1

expression was predominantly observed in metastatic specimens. In the present study, no expression of PD-L1 was observed in any patient using immunohistochemistry. Moreover, PD-L1 mRNA expression levels were extremely low in both the SS cell lines and in the clinical specimens. The data therefore suggest that SS escapes from the immune system by a mechanism other than the PD-1/PD-L1 axis. The role of PD-L2 in the tumor microenvironment is less understood compared to that of PD-L1 in sarcomas. Zheng et al³⁶ first examined the expression of PD-L2 in 234 patients with musculoskeletal tumors including 127 cases of patients with SS, and 26 cases (20.5%) of SS were positive for PD-L2 by immunohistochemistry. However, the prognostic impact of PD-L1 and PD-L2 in SS remains unclear and further experiments are required to show the mechanism underlying the association among PD-L1, PD-L2, and prognosis.

Macrophages represent a major component of tumor immune cells, and may be classified into tumoricidal M1-like macrophages or pro-tumoral M2-like macrophages.³⁷ M2-like macrophages may be involved in tumor progression by promoting angiogenesis, suppressive adaptive immunity, tumor cell proliferation, and metastasis.^{37,38} CD163 is considered a marker for M2-like macrophages and a high number of infiltrating CD163+ macrophages represents a poor prognostic predictor in several cancers.³⁸⁻⁴⁰ However, the prognostic impact of CD163+ macrophages in patients with SS is still unknown. To our knowledge, this is the first report that has examined the association between the infiltration of CD163+ macrophages and SS prognosis in a patient cohort. In the present study, we showed that a higher infiltration of CD163+ macrophages was associated with poor prognosis and the progression of SS. This suggests that CD163+ macrophages might play a key role in the progression of the disease in patients with SS, more so than PD-L1. Indeed, although a case report discussed the efficacy of PD-1 blockade therapy in advanced sarcoma,⁴¹ the efficacy of a therapy targeting PD-1 in soft tissue sarcoma is limited.⁴² Toulmonde et al reported that CD163+ macrophages were markedly infiltrated, whereas PD-L1 expression was low in most advanced soft tissue sarcomas. Therefore, immunosuppression in soft tissue sarcomas may be as a result of the action of CD163+ macrophages and may represent the reason why the efficacy of PD-1-targeting therapy is limited.⁴² Moreover, Shiraishi et al⁴³ indicated that CD163 is related to protumoral activation of macrophages, and development and progression of tumor by interleukin-6 from macrophages. Meanwhile, it was reported that tumor-associated macrophages, especially CD163+ CD204+, expressed higher levels of PD-L1 than other macrophage subsets.^{44,45} Furthermore, PD-L1 expression on dendritic cells and macrophages in the tumor microenvironment is important for immunosuppression and resistance to PD-1/PD-L1 blockade therapy.^{44,46} In the present study, there was no correlation between the number of CD163+ macrophages and PD-L1 or PD-L2 mRNA expression. This discrepancy might be due to the limited number of specimens available for real-time PCR. Further analyses with large samples are needed to explain the association between CD163+ macrophages and PD-L1/PD-L2 expression and the clinical significance of CD163+ macrophages in patients with SS.

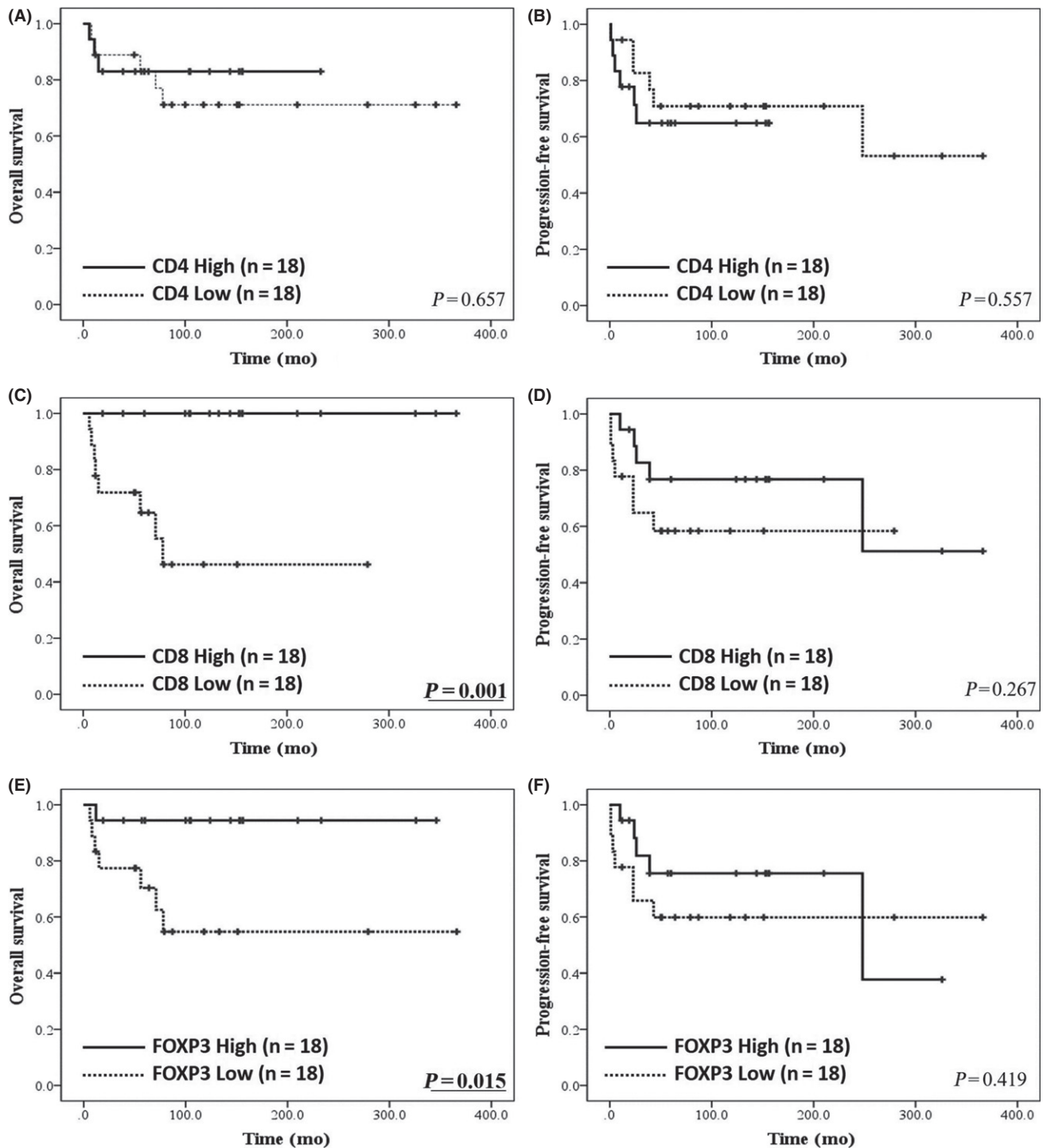


FIGURE 5 Kaplan-Meier curves illustrating overall survival (OS) and progression-free survival (PFS) for CD4, CD8, FOXP3, CD163, human leukocyte antigen (HLA) class I, and programmed death ligand 1 (PD-L1)/programmed death ligand 2 (PD-L2) mRNA expression. Survival differences were analyzed using the log-rank test. The increased infiltration of CD4+ lymphocytes was not associated with OS or PFS (A,B). Patients with a higher infiltration of CD8+ or FOXP3+ lymphocytes showed a significant improvement in OS, but not PFS (C-F). There was a significant difference for both OS and PFS in patients showing a lower infiltration of CD163+ macrophages (G,H). Expression of HLA class I and that of PD-L1 and PD-L2 mRNA were not significantly associated with either OS or PFS (I-N)

Another mechanism for resistance to PD-1 targeted therapy was reported, showing that macrophages arrested CD8+ lymphocytes by shaping long-lasting interaction and inhibiting CD8+ lymphocyte migration into tumor nests and contact with tumor cells. Depletion

of macrophages allowed CD8+ lymphocytes to migrate into tumor nests and scan tumor cells to improve the efficacy of PD-1 targeting immunotherapy.⁴⁷ In this study, although the density of CD8+ lymphocytes was correlated with that of CD163+ macrophages, each

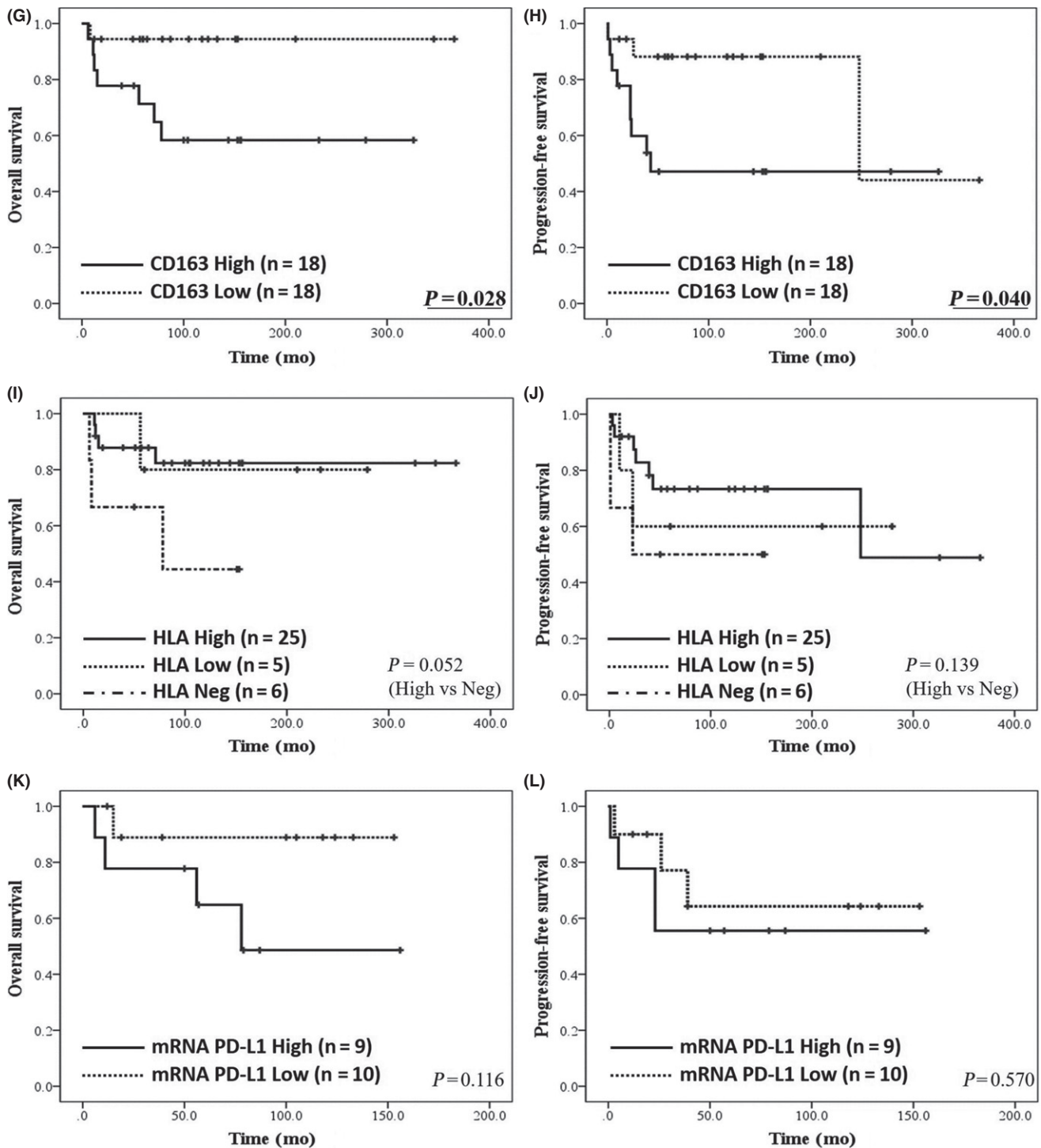


FIGURE 5 (Continued)

factor had an opposite impact on prognosis. This may have resulted from the fact that it is difficult to definitively distinguish tumor nests from stroma in SS, unlike other types of cancer. Further studies are needed to verify the interaction between CD8+ lymphocytes and CD163+ macrophages in patients with SS. Taken together, it is suggested that macrophage-targeting therapy or combining approaches targeting macrophages and PD-1 may represent a promising immunotherapy in patients with SS.

The present study has several limitations. First, as this was a retrospective study, the patient's baseline characteristics varied. Thus, this study comprises patients with distant metastasis at diagnosis, and this might have influenced our results. To confirm these results, further studies are needed to evaluate a greater number of tumor specimens with a similar presentation status. Moreover, the evaluation of metastatic specimens was beyond the scope of our study, and we could not address the tumor immune microenvironment in metastatic tumors.

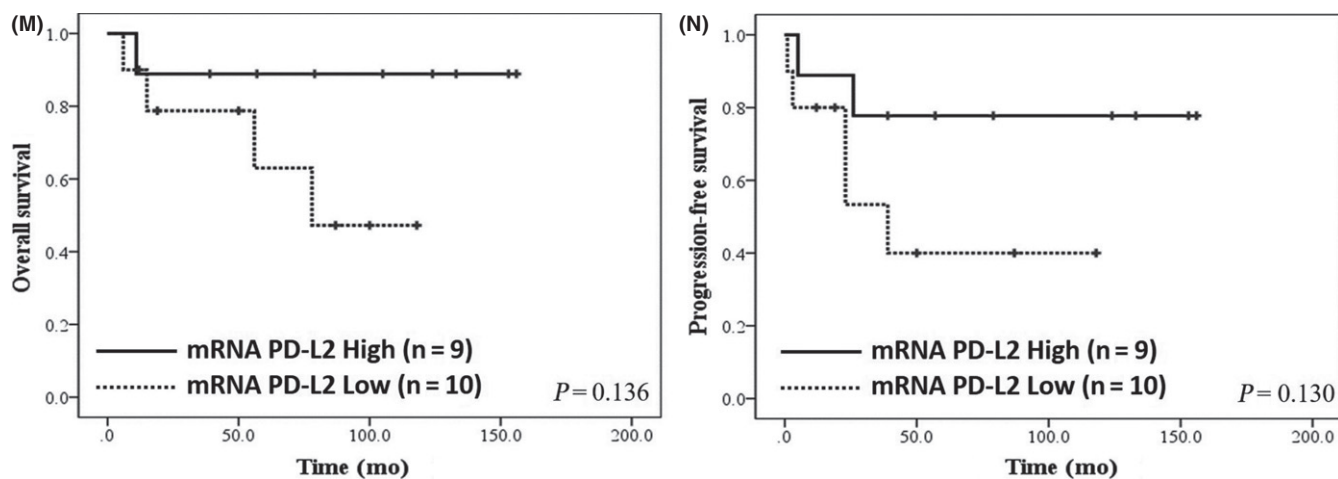


FIGURE 5 (Continued)

Finally, we evaluated the expression of PD-L1 using one antibody only. Therefore, our results could be influenced by the antibody used. To remedy this problem, a standardized method for testing the expression of PD-L1 using immunohistochemistry is required.

In conclusion, we showed that PD-L1 was not expressed in any of the patients, as assessed by immunohistochemistry, and that the PD-

L1 mRNA expression level was low in the tumor microenvironment of patients with SS. In addition, our results showed that a lower infiltration of CD8+ and FOXP3+ lymphocytes and a higher infiltration of CD163+ macrophages were associated with an unfavorable prognosis in patients with SS. We also showed that the infiltration of CD163+ macrophages might play an important role in disease progression in patients with SS. We believe these data offer novel insights into the SS tumor immune microenvironment and will provide a foundation for developing immunotherapies for patients with SS.

TABLE 3 Univariate analysis of prognostic clinicopathological factors for OS and PFS

Variable	Category	No. of patients	P-value	
			OS	PFS
Age at diagnosis, y	<20 (vs ≥20)	6	0.553	0.970
Gender	Male (vs female)	23	0.500	0.487
Location	Extremity (vs trunk)	33	0.758	0.194
Size, cm	>5 (vs ≤5)	14	0.094	0.570
Translocation	SYT-SSX 1 (vs SYT-SSX 2)	24	0.734	0.213
Stage (AJCC system)	Stage II (vs stage III)	14	0.310	0.874
	Stage III (vs stage IV)	19	<0.001	<0.001
Surgical margin	Adequate (vs inadequate)	30	0.981	0.441

AJCC, American Joint Committee on Cancer systems 7th edition staging manual; OS, overall survival; PFS, progression-free survival.

TABLE 4 Multivariate analysis for clinicopathological and immunohistochemical parameters

Variables	P-value	HR	95% CI
Stage (AJCC system)	0.010	15.371	1.930-122.454
No. of CD8+ lymphocytes	0.030	0.781	0.625-0.977
No. of CD163+ macrophages	0.037	1.078	1.005-1.157

AJCC, American Joint Committee on Cancer systems 7th edition staging manual; CI, confidence interval; HR, hazard ratio.

ACKNOWLEDGMENTS

The authors thank Ms K. Tanaka and Mr Y. Tanaka (Division of Orthopedic Surgery, Niigata University Graduate School of Medical and Dental Sciences, Niigata, Japan) for their technical assistance. The authors are also grateful to the staff in the Departments of Pathology of the Niigata University Medical and Dental Hospital and the Niigata Cancer Center Hospital for their technical assistance.

CONFLICTS OF INTEREST

Authors declare no conflicts of interest for this article.

ORCID

Naoki Oike  <http://orcid.org/0000-0003-3067-392X>

REFERENCES

- Brennan B, Stevens M, Kelsey A, Stiller CA. Synovial sarcoma in childhood and adolescence: a retrospective series of 77 patients registered by the Children's Cancer and Leukaemia Group between 1991 and 2006. *Pediatr Blood Cancer*. 2010;55:85-90.
- Italiano A, Penel N, Robin YM, et al. Neo/adjuvant chemotherapy does not improve outcome in resected primary synovial sarcoma: a study of the French Sarcoma Group. *Ann Oncol*. 2009;20:425-430.

3. Lewis JJ, Antonescu CR, Leung DH, et al. Synovial sarcoma: a multivariate analysis of prognostic factors in 112 patients with primary localized tumors of the extremity. *J Clin Oncol.* 2000;18:2087-2094.
4. Brodin B, Haslam K, Yang K, et al. Cloning and characterization of spliced fusion transcript variants of synovial sarcoma: SYT/SSX4, SYT/SSX4v, and SYT/SSX2v. Possible regulatory role of the fusion gene product in wild type SYT expression. *Gene.* 2001;268:173-182.
5. Kawai A, Woodruff J, Healey JH, Brennan MF, Antonescu CR, Ladanyi M. SYT-SSX gene fusion as a determinant of morphology and prognosis in synovial sarcoma. *N Engl J Med.* 1998;338:153-160.
6. Jungbluth AA, Antonescu CR, Busam KJ, et al. Monophasic and biphasic synovial sarcomas abundantly express cancer/testis antigen NY-ESO-1 but not MAGE-A1 or CT7. *Int J Cancer.* 2001;94:252-256.
7. Lai JP, Robbins PF, Raffeld M, et al. NY-ESO-1 expression in synovial sarcoma and other mesenchymal tumors: significance for NY-ESO-1-based targeted therapy and differential diagnosis. *Mod Pathol.* 2012;25:854-858.
8. Kawaguchi S, Tsukahara T, Ida K, et al. SYT-SSX breakpoint peptide vaccines in patients with synovial sarcoma: a study from the Japanese Musculoskeletal Oncology Group. *Cancer Sci.* 2012;103:1625-1630.
9. Dutoit V, Taub RN, Papadopoulos KP, et al. Multiepitope CD8(+) T cell response to a NY-ESO-1 peptide vaccine results in imprecise tumor targeting. *J Clin Invest.* 2002;110:1813-1822.
10. Robbins PF, Kassim SH, Tran TL, et al. A pilot trial using lymphocytes genetically engineered with an NY-ESO-1-reactive T-cell receptor: long-term follow-up and correlates with response. *Clin Cancer Res.* 2015;21:1019-1027.
11. Herbst RS, Baas P, Kim D-W, et al. Pembrolizumab versus docetaxel for previously treated, PD-L1-positive, advanced non-small-cell lung cancer (KEYNOTE-010): a randomised controlled trial. *Lancet.* 2016;387:1540-1550.
12. Robert C, Schachter J, Long GV, et al. Pembrolizumab versus ipilimumab in advanced melanoma. *N Engl J Med.* 2015;372:2521-2532.
13. Armand P, Shipp MA, Ribrag V, et al. Programmed death-1 blockade with Pembrolizumab in patients with classical Hodgkin Lymphoma after Brentuximab Vedotin failure. *J Clin Oncol.* 2016;34:3733-3739.
14. Tanaka K, Miyata H, Sugimura K, et al. Negative influence of programmed death-1-ligands on the survival of esophageal cancer patients treated with chemotherapy. *Cancer Sci.* 2016;107:726-733.
15. Muenst S, Schaerli A, Gao F, et al. Expression of programmed death ligand 1 (PD-L1) is associated with poor prognosis in human breast cancer. *Breast Cancer Res Treat.* 2014;146:15-24.
16. Lipson EJ, Vincent JG, Loyo M, et al. PD-L1 expression in the Merkel cell carcinoma microenvironment: association with inflammation, Merkel cell polyomavirus, and overall survival. *Cancer Immunol Res.* 2013;1:54-63.
17. Yearley JH, Gibson C, Yu N, et al. PD-L2 expression in human tumors: relevance to anti-PD-1 therapy in cancer. *Clin Cancer Res.* 2017;23:3158-3167.
18. van Erp AEM, Versleijen-Jonkers YMH, Hillebrandt-Roeffen MHS, et al. Expression and clinical association of programmed cell death-1, programmed death-ligand-1 and CD8(+) lymphocytes in primary sarcomas is subtype dependent. *Oncotarget.* 2017;8:71371-71384.
19. Nowicki TS, Akiyama R, Huang RR, et al. Infiltration of CD8 T cells and expression of PD-1 and PD-L1 in synovial sarcoma. *Cancer Immunol Res.* 2017;5:118-126.
20. Kim C, Kim EK, Jung H, et al. Prognostic implications of PD-L1 expression in patients with soft tissue sarcoma. *BMC Cancer.* 2016;16:434.
21. D'Angelo SP, Shoushtari AN, Agaram NP, et al. Prevalence of tumor-infiltrating lymphocytes and PD-L1 expression in the soft tissue sarcoma microenvironment. *Hum Pathol.* 2015;46:357-365.
22. Edge SB, Compton CC. The American Joint Committee on Cancer: the 7th edition of the AJCC cancer staging manual and the future of TNM. *Ann Surg Oncol.* 2010;17:1471-1474.
23. Koirala P, Roth ME, Gill J, et al. Immune infiltration and PD-L1 expression in the tumor microenvironment are prognostic in osteosarcoma. *Sci Rep.* 2016;6:30093.
24. Sonobe H, Manabe Y, Furihata M, et al. Establishment and characterization of a new human synovial sarcoma cell line, HS-SY-II. *Lab Invest.* 1992;67:498-505.
25. Kawai A, Naito N, Yoshida A, et al. Establishment and characterization of a biphasic synovial sarcoma cell line, SYO-1. *Cancer Lett.* 2004;204:105-113.
26. Li G, Kawashima H, Ji L, et al. Frequent absence of tumor suppressor FUS1 protein expression in human bone and soft tissue sarcomas. *Anticancer Res.* 2011;31:11-21.
27. Boxberg M, Steiger K, Lenze U, et al. PD-L1 and PD-1 and characterization of tumor-infiltrating lymphocytes in high grade sarcomas of soft tissue - prognostic implications and rationale for immunotherapy. *Oncoimmunology.* 2018;7:e1389366.
28. Gooden MJ, de Bock GH, Leffers N, Daemen T, Nijman HW. The prognostic influence of tumour-infiltrating lymphocytes in cancer: a systematic review with meta-analysis. *Br J Cancer.* 2011;105:93-103.
29. Fujii H, Arakawa A, Utsumi D, et al. CD8(+) tumor-infiltrating lymphocytes at primary sites as a possible prognostic factor of cutaneous angiosarcoma. *Int J Cancer.* 2014;134:2393-2402.
30. Yabe H, Tsukahara T, Kawaguchi S, et al. Prognostic significance of HLA class I expression in Ewing's sarcoma family of tumors. *J Surg Oncol.* 2011;103:380-385.
31. Sakaguchi S. Naturally arising Foxp3-expressing CD25+CD4+ regulatory T cells in immunological tolerance to self and non-self. *Nat Immunol.* 2005;6:345-352.
32. Wolf D, Wolf AM, Rumpold H, et al. The expression of the regulatory T cell-specific forkhead box transcription factor FoxP3 is associated with poor prognosis in ovarian cancer. *Clin Cancer Res.* 2005;11:8326-8331.
33. Que Y, Xiao W, Guan Y-X, et al. PD-L1 expression is associated with FOXP3+ regulatory T-cell infiltration of soft tissue sarcoma and poor patient prognosis. *J Cancer.* 2017;8:2018-2025.
34. Miyara M, Yoshioka Y, Kitoh A, et al. Functional delineation and differentiation dynamics of human CD4+ T cells expressing the FoxP3 transcription factor. *Immunity.* 2009;30:899-911.
35. Kim JR, Moon YJ, Kwon KS, et al. Tumor infiltrating PD1-positive lymphocytes and the expression of PD-L1 predict poor prognosis of soft tissue sarcomas. *PLoS ONE.* 2013;8:e82870.
36. Zheng B, Ren T, Huang Y, et al. PD-1 axis expression in musculoskeletal tumors and antitumor effect of nivolumab in osteosarcoma model of humanized mouse. *J Hematol Oncol.* 2018;11:16.
37. Mantovani A, Sozzani S, Locati M, Allavena P, Sica A. Macrophage polarization: tumor-associated macrophages as a paradigm for polarized M2 mononuclear phagocytes. *Trends Immunol.* 2002;23:549-555.
38. Takeya M, Komohara Y. Role of tumor-associated macrophages in human malignancies: friend or foe? *Pathol Int.* 2016;66:491-505.
39. Hasita H, Komohara Y, Okabe H, et al. Significance of alternatively activated macrophages in patients with intrahepatic cholangiocarcinoma. *Cancer Sci.* 2010;101:1913-1919.
40. Medrek C, Pontén F, Jirström K, Leandersson K. The presence of tumor associated macrophages in tumor stroma as a prognostic marker for breast cancer patients. *BMC Cancer.* 2012;12:306.
41. McCaughan GJ, Fulham MJ, Mahar A, et al. Programmed cell death-1 blockade in recurrent disseminated Ewing sarcoma. *J Hematol Oncol.* 2016;9:48.
42. Toulmonde M, Penel N, Adam J, et al. Use of PD-1 targeting, macrophage infiltration, and IDO pathway activation in sarcomas: a phase 2 clinical trial. *JAMA Oncol.* 2018;4:93-97.
43. Shiraishi D, Fujiwara Y, Horlad H, et al. CD163 is required for protumoral activation of macrophages in human and murine sarcoma. *Cancer Res.* 2018;78:3255-3266.

44. Horlad H, Ma C, Yano H, et al. An IL-27/Stat3 axis induces expression of programmed cell death 1 ligands (PD-L1/2) on infiltrating macrophages in lymphoma. *Cancer Sci*. 2016;107:1696-1704.
45. Kubota K, Moriyama M, Furukawa S, et al. CD163+ CD204+ tumor-associated macrophages contribute to T cell regulation via interleukin-10 and PD-L1 production in oral squamous cell carcinoma. *Sci Rep*. 2017;7:1755.
46. Lin H, Wei S, Hurt EM, et al. Host expression of PD-L1 determines efficacy of PD-L1 pathway blockade-mediated tumor regression. *J Clin Invest*. 2018;128:805-815.
47. Peranzoni E, Lemoine J, Vimeux L, et al. Macrophages impede CD8 T cells from reaching tumor cells and limit the efficacy of

anti-PD-1 treatment. *Proc Natl Acad Sci USA*. 2018;115:E4041-E4050.

How to cite this article: Oike N, Kawashima H, Ogose A, et al. Prognostic impact of the tumor immune microenvironment in synovial sarcoma. *Cancer Sci*. 2018;109:3043-3054. <https://doi.org/10.1111/cas.13769>