

Serum levels of soluble programmed cell death ligand 1 as a prognostic factor on the first-line treatment of metastatic or recurrent gastric cancer

Naoki Takahashi¹ · Satoru Iwasa¹ · Yusuke Sasaki¹ · Hirokazu Shoji¹ · Yoshitaka Honma¹ · Atsuo Takashima¹ · Natsuko Tsuda Okita¹ · Ken Kato¹ · Tetsuya Hamaguchi¹ · Yasuhide Yamada¹

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Abstract

Purpose Immune checkpoint molecules are key targets for the treatment of various malignancies. Due to the heterogeneity of advanced gastric cancer (GC), the role of programmed cell death ligand 1 (PD-L1) expression as a tumor biomarker remains controversial. In this study, the prognostic value of soluble PD-L1 (sPD-L1) levels in serum samples was assessed in patients with metastatic GC.

Methods All patients received first-line treatment with fluoropyrimidine and platinum chemotherapy, and trastuzumab was added for HER2-positive patients. Serum levels of sPD-L1 were measured by enzyme-linked immunosorbent assay.

Results Among 75 metastatic GC patients, the median serum sPD-L1 level was 0.704 ng/ml (range <0.156–3.214). Serum sPD-L1 was significantly higher in patients with a high versus a low white blood cell count at baseline. When the cutoff value was set as the median, multivariate analyses showed that high sPD-L1 levels were associated with worse overall survival compared with low sPD-L1 levels (HR 2.218, 95 % CI 1.139–4.320, $P = 0.019$). Regardless of HER2 status, overall survival tended to be shorter in patients with high sPD-L1 compared with low sPD-L1.

There was no significant association between sPD-L1 level and progression-free survival on the first-line treatment of metastatic GC.

Conclusions High serum levels of sPD-L1 correlated with worse overall survival on the first-line chemotherapy in metastatic GC patients.

Keywords Gastric cancer · Prognostic factor · PD-L1 · Serum

Introduction

Gastric cancer (GC) is prevalent in males and females, being the third and fifth leading cause of cancer-related mortality in the two respective genders worldwide (Torre et al. 2012). The incidence of GC is highest in Eastern Asia, Central and Eastern Europe, and South America and lowest in North America and most African countries. Although endoscopic or curative resection may prolong survival in patients with early GC, the prognosis for metastatic or recurrent GC remains poor, despite the development of novel molecularly targeted drugs (Bang et al. 2010; Fuchs et al. 2014; Wilke et al. 2014).

The tumor microenvironment consists of stem, stromal, and endothelial cells, and a variety of immune cells (Hanahan and Coussens 2012). Programmed cell death 1 (PD-1) and its ligand (PD-L1) are members of the B7 family, which negatively regulates the immune system and plays important roles in maintaining self-tolerance, preventing autoimmunity, and protecting against immune collateral damage in normal tissue (Murdoch et al. 2008; Yang et al. 2008; Joyce and Pollard 2009). PD-L1 is a transmembrane protein that is often overexpressed in malignant tumors, and inhibits T cell-mediated immune attack through binding to

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✉ Naoki Takahashi
naoki19800623@gmail.com

Yasuhide Yamada
yasuhide-yamada@amed.go.jp

¹ Gastrointestinal Medical Oncology Division, National Cancer Center Hospital, 5-1-1 Tsukiji, Chuo-ku, Tokyo 104-0045, Japan

PD-1 on tumor-specific T cells. PD-L1 expression is regulated by several intracellular mechanisms, including activation of the STAT3 pathway by inflammatory cytokines such as interferon gamma (IFN- γ), as well as the mitogen-activated protein kinase (MAPK)/nuclear factor-kappa B (NF- κ B) pathway and the phosphatidylinositide 3-kinase (PI3K)/AKT pathway (Qian et al. 2008; Crane et al. 2009). PD-L1- and PD-1-targeted therapies have been developed for several cancer types. The efficacy and safety profile of an anti-PD-1 antibody has been confirmed in randomized phase 3 trials in malignant melanoma, non-small cell lung cancer, and renal cell carcinoma (Borghaei et al. 2015; Motzer et al. 2015; Robert et al. 2015).

Previous studies indicate that PD-L1 expression is a poor prognostic factor in several histological types of malignant tumor (Thompson et al. 2006; Nomi et al. 2007; Kim et al. 2013; Muenst et al. 2014; Huang et al. 2015; Paydas et al. 2015; Wang et al. 2015a, b; Wu et al. 2015; Cierna et al. 2016). Although overexpression of PD-L1 in tumor tissue, assessed by immunohistochemistry (IHC), has been linked to responses to anti-PD-L1 antibody treatment, the value of PD-L1 as a prognostic or predictive biomarker remains controversial. This may be explained by the heterogeneity of PD-L1 expression in tumor tissue, and inconsistencies in the definition of PD-L1 positivity and diagnostic antibodies (Callea et al. 2015; Madore et al. 2015; McLaughlin et al. 2016). In addition, PD-L1 expression in tumor tissue is affected by the timing of biopsy during treatment, the composition of tumor tissue, and differences between biopsies from the primary lesion and metastatic sites (Kerr et al. 2015).

PD-L1 expression can be detected on the surface of tumor and immune cells by IHC, and in blood samples by enzyme-linked immunosorbent assay (ELISA). PD-L1 is a component of normal human serum, and soluble PD-L1 (sPD-L1) is released from PD-L1-positive cell lines (Chen et al. 2011). Serum sPD-L1 levels are higher in patients with malignant cancer than in healthy individuals, and high sPD-L1 was a poor prognostic factor during standard treatment of hematopoietic malignancies in recent studies (Rossille et al. 2014; Wang et al. 2015a, b). To date, few investigations have evaluated the relationship between serum sPD-L1 levels and the prognosis of metastatic GC patients treated with systemic chemotherapy, and the present study was conducted to address this issue.

Materials and methods

Patients and sample collection

Patients with histologically diagnosed gastric adenocarcinoma and radiologic confirmation of metastatic or recurrent

lesions were enrolled. Other inclusion criteria were as follows: age ≥ 20 years, Eastern Cooperative Oncology Group (ECOG) Performance Status (PS) 0–2, adequate hematological and organ function, at least one measurable target lesion, no prior chemotherapy or radiotherapy, first-line cancer treatment, no brain or central nervous system metastases, no coexisting cancers except for gastroesophageal junction adenocarcinoma, no uncontrolled complications, sufficient serum samples at baseline, and the patient's written informed consent for use of clinical data and materials.

Patients' sera were obtained from residual blood samples, which were collected for other laboratory tests, between September 2011 and July 2015, within 2 weeks of initiating first-line treatment. Separated serum samples were immediately stored at -20°C in the Biobank of the National Cancer Center Hospital, Tokyo, Japan, until analysis. In advanced GC, human endothelial growth factor receptor 2 (HER2) positivity tends to be less frequent than HER2 negativity. To enable the prognostic value of sPD-L1 levels to be evaluated in both HER2-positive and HER2-negative patients, more HER2-positive patients were enrolled in this study, regardless of the outcome of chemotherapy.

HER2 expression was determined by IHC (HerceptTest, Dako, Denmark) of formalin-fixed paraffin-embedded tumor specimens from all patients. HER2 staining was graded on a scale of 0–3 as follows: 0 = no reactivity or membranous reactivity in $\geq 10\%$ of cells; 1+ = faint/barely perceptible membranous reactivity in $\geq 10\%$ of cells or reactivity in a portion of the cell membrane; 2+ = weak-to-moderate complete or basolateral membranous reactivity in $\geq 10\%$ of cells; and 3+ = strong complete or basolateral membranous reactivity in $\geq 10\%$ of cells. In tumor specimens with IHC scores of 2+, dual in situ hybridization (DISH; Ventana INFORM HER2 Dual Color ISH, Roche) was also performed. HER2 IHC3+ and IHC2+/DISH-positive (HER2/CEP17 ratio ≥ 2) specimens were defined as HER2 positive, and other specimens were defined as HER2 negative.

Informed consent of Biobank was obtained from all individual participants included in the study. The study protocol was approved by the local institutional review board.

Treatment

All patients received first-line chemotherapy comprising a fluoropyrimidine (S-1, capecitabine or 5-fluorouracil) and platinum (cisplatin or oxaliplatin). HER2-negative patients received either dual therapy consisting of S-1 plus cisplatin (CS), S-1 plus oxaliplatin (SOX), or capecitabine plus oxaliplatin (XELOX), or triple therapy comprising docetaxel plus CS (DCS). HER2-positive patients received triple therapy comprising trastuzumab with either

5-fluorouracil plus cisplatin (FP) or capecitabine plus cisplatin (XP). The chemotherapeutic regimens, which are summarized in Supplementary material 1A, were based on previous clinical trials (Bang et al. 2010; Koizumi et al. 2008, 2012; Kim et al. 2012; Yamada et al. 2015). Patients continued to receive treatment until disease progression or intolerable toxicity from chemotherapy. In the case of severe adverse events, dose modifications were made according to the physician's judgment. Responses to treatment were evaluated every 2–3 months by contrast-enhanced computed tomography (CT).

ELISA

Serum PD-L1 levels were measured using a commercially available ELISA kit (USCN, Wuhan, China). ELISA was conducted as follows: (1) all reagents, standard dilutions, and samples were prepared according to the manufacturer's instructions; (2) 100 μ L of the standard and sample were added to each well; (3) plates were covered with a plate sealer and incubated at 37 °C for 2 h; (4) after aspiration, 100 μ L of detection reagent A (1:100) was added to each well, and the plates were re-sealed and incubated at 37 °C for 2 h; (5) each well was aspirated and washed four times in wash buffer; (6) 100 μ L of detection reagent B (1:100) was added to each well, and the plates were re-sealed and incubated for 0.5 h at 37 °C; (7) each well was aspirated and washed five times, 90 μ L of substrate solution was added, and the plates were newly sealed and incubated in a dark room for 20 min at 37 °C; (8) 50 μ L of stop solution was added to each well, and absorbance at 450 nm was immediately measured with a plate reader.

Disease assessment and statistical analyses

Therapeutic responses were categorized as complete response (CR), partial response (PR), stable disease (SD), disease progression (PD), and non-evaluable (NE), according to the Response Evaluation Criteria in Solid Tumors (RECIST) criteria version 1.1. The overall response rate (ORR) was defined as the proportion of the study population with a best response of CR or PR. The disease control rate (DCR) was defined as the proportion of patients with a best response of CR, PR, or SD. Progression-free survival (PFS) was defined as the time from initiation of first-line treatment to disease progression or death without evidence of progression. Overall survival (OS) was defined as the time from initiation of first-line treatment to death or last follow-up. Survival curves for PFS and OS were estimated by the Kaplan–Meier method, and differences were evaluated by the log-rank test. Hazard ratios (HRs) and 95 % confidence intervals (CIs) were estimated with a Cox proportional hazards model. HRs for univariate analyses were

calculated using the covariates of age (younger vs. older, cutoff; median), PS (0–1 vs. 2), gender (male vs. female), histological type (intestinal vs. distal and mixed), stage (stage IV vs. recurrence), HER2 status (positive vs. negative), prior gastrectomy (yes vs. no), number of metastatic sites (1 vs. ≥ 2), peritoneal dissemination (yes vs. no), liver metastasis (yes vs. no), chemotherapeutic regimen (dual vs. triple), and serum sPD-L1 levels (high vs. low, cutoff level: median). HRs in multivariate analyses were adjusted by first-line treatment, serum levels of PD-L1, and patient characteristics as covariates, all of which had a *P*-value of < 0.2 in univariate analysis.

Differences in the distribution of two variables were evaluated using Fisher's exact test or χ^2 test, as appropriate. Differences in the distribution of more than two variables were evaluated by the Kruskal–Wallis test. Receiver operating characteristic (ROC) curves were generated to determine the optimal cutoff value for sPD-L1 levels as a continuous variable. All tests were two-sided and *P* < 0.05 was considered to denote statistical significance. All statistical analyses were performed using SPSS software, version 19 (IBM, Tokyo, Japan).

Results

The baseline characteristics of the study population, which comprised 75 patients, are summarized in Table 1. The median age was 67.0 years (range: 39–79 years), and 96.0 % were ECOG PS 0–1. Histologically, tumors were diffuse (54.7 %), intestinal (38.7 %), and mixed type (6.6 %). Analysis of HER2 status revealed that 45 (60.0 %) and 30 (40.0 %) patients were HER2 negative and HER2 positive, respectively. Among all 75 patients, 67 patients (89.3 %) received at least one subsequent treatment after failure of first-line treatment. As shown in Supplementary material 1B, subsequent treatments were a taxane (89.6 %), irinotecan (49.3 %), ramucirumab (6.0 %), and other drugs (20.9 %). The frequency of ramucirumab administration was low because of its recent approval in Japan in March 2015.

Serum sPD-L1 levels

As shown in Fig. 1, the median serum sPD-L1 value of the study population was 0.704 ng/ml (range: < 0.156 –3.214 ng/ml). In 29 patients, serum sPD-L1 levels were lower than the limit of detection. The distribution of serum sPD-L1 levels by response to first-line treatment was evaluated. In patients with a PR, SD, and PD, median serum sPD-L1 levels were 0.707, 0.622, and 0.751 ng/ml, respectively. There was no significant difference in sPD-L1 levels according to the response to first-line treatment.

Table 1 Patient characteristics

Age, years	67.0 (39–79)
<i>Gender</i>	
Male	58 (77.3)
Female	17 (22.7)
<i>ECOG PS</i>	
0–1	72 (96.0)
2	3 (4.0)
<i>Histological type</i>	
Intestinal	29 (38.7)
Diffuse	41 (54.7)
Mixed	5 (6.6)
<i>Stage</i>	
Stage IV	62 (82.7)
Recurrence	13 (17.3)
<i>HER2 status</i>	
Negative	45 (60.0)
Positive	30 (40.0)
<i>Number of metastatic sites</i>	
0–1	40 (53.3)
2≤	35 (46.7)
<i>Metastatic organs</i>	
Lymph nodes	56 (74.7)
Liver	34 (45.3)
Lung	8 (10.7)
Peritoneum	24 (32.0)
<i>Chemotherapeutic regimen</i>	
Dual	65 (86.7)
Triplet	10 (13.3)
<i>Subsequent treatment</i>	
Yes	67 (89.3)
No	8 (10.7)

All values are expressed as the number (%) of patients, except for age, which is expressed as the median (range)

ECOG PS Eastern Cooperative Oncology Group Performance Status, *Dual* fluoropyrimidine/platinum, *Triplet* docetaxel/fluoropyrimidine/cisplatin

The association between sPD-L1 levels and inflammatory markers, including white cell count (WBC), C-reactive protein (CRP), albumin, and hemoglobin, was explored. As shown in Fig. 2, patients with a high WBC (cutoff; 8000/ μ L) had higher serum sPD-L1 levels than those with a low WBC ($P = 0.043$). There were no correlations between serum sPD-L1 levels and other inflammatory markers.

Prognostic analyses of PFS and OS on the first-line treatment

The cutoff value for serum sPD-L1 was initially set as the median. Univariate and multivariate analyses of

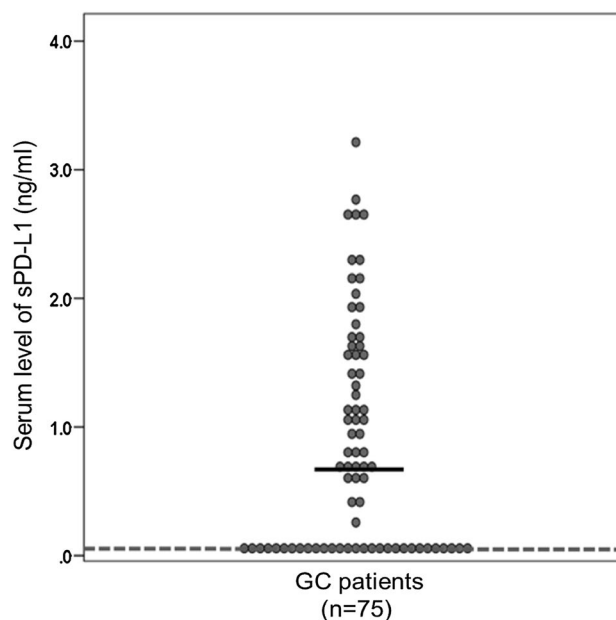


Fig. 1 Serum levels of sPD-L1 in patients with metastatic or recurrent gastric cancer. The distribution of sPD-L1 levels in serum (ng/ml) is shown (A). The *dashed line* represents the minimum detectable level (0.157 ng/ml). The *solid bar* represents the median sPD-L1 level of all gastric cancer patients

PFS are shown in Supplementary material 2. Univariate analysis showed a trend toward high sPD-L1 levels being associated with low PFS on the first-line treatment (HR 1.580, 95 % CI 0.958–2.605, $P = 0.073$). However, multivariate analysis demonstrated that serum sPD-L1 was not independently related to PFS (HR 1.261, 95 % CI 0.711–2.236, $P = 0.427$). Univariate and multivariate analyses of OS are given in Table 2. Univariate analysis identified stage IV (HR 2.237, 95 % CI 1.003–4.990, $P = 0.049$), HER2 positive (HR 0.537, 95 % CI 0.293–0.983, $P = 0.044$), and high sPD-L1 (HR 2.647, 95 % CI 1.476–4.745, $P = 0.001$). Multivariate analysis indicated that high serum sPD-L1 was independently associated with low OS (HR 2.218, 95 % CI 1.139–4.320, $P = 0.019$).

The baseline characteristics of the high and low sPD-L1 subgroups are shown in Supplementary material 3. No significant differences were found in the baseline characteristics of patients with high as opposed to low serum sPD-L1 levels, but stage IV was more frequently observed in patients with high levels of sPD-L1 compared with those with low levels of sPD-L1. The frequency of subsequent treatment after failure of first-line treatment was not different between the high and low sPD-L1 groups (Supplementary material 1B). The frequency of administration of each type of drug such as taxanes and irinotecan was similar between groups.

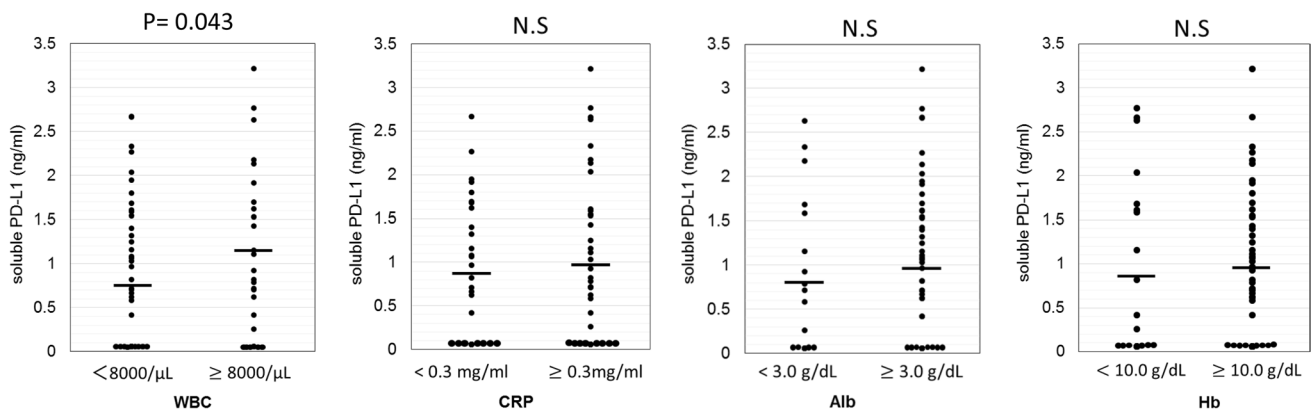


Fig. 2 Association between serum sPD-L1 levels and inflammatory factors. The inflammatory factors evaluated were white cell count (WBC), C-reactive protein (CRP), albumin (Alb), and hemoglobin (Hb). *NS* not significant

PFS and OS curves using median sPD-L1 as the cutoff value

Figure 3 shows PFS and OS curves using median sPD-L1 as the cutoff value. Median PFS for high and low sPD-L1 was 5.2 months (95 % CI 3.7–7.5) and 7.4 months (95 % CI 4.6–10.3), respectively. There was no significant difference in PFS, but patients with high sPD-L1 tended to have shorter PFS than those with low sPD-L1 levels (log-rank test, $P = 0.070$). Median OS for high and low sPD-L1 was 13.2 months (95 % CI 11.1–16.7) and 21.6 months (95 % CI 11.1–35.3), respectively. Patients with high sPD-L1 levels had shorter OS than those with low sPD-L1 s (log-rank test, $P = 0.0009$).

The difference between survival curves in HER2-negative and HER2-positive patients was evaluated, as illustrated in Fig. 4. In HER2-negative patients, median PFS for high and low sPD-L1 was not different (4.9 vs. 6.5 months, $P = 0.562$), while median OS tended to be shorter in those with high sPD-L1 (14.0 vs 21.6 months, $P = 0.076$). In HER2-positive patients, median PFS was similar for high and low sPD-L1 levels (7.8 vs. 7.9 months, $P = 0.379$), whereas median OS was shorter for high sPD-L1 compared with low sPD-L1 (13.2 months vs. not reached, $P = 0.0196$).

Survival curves using the optimal cutoff value for sPD-L1

The optimal cutoff value of sPD-L1 was estimated by ROC curve analysis to set at that for the best DCR. As a result, the cutoff value of sPD-L1 was 1.081 ng/ml. Survival curves for PFS and OS using the optimal cutoff value are shown in Fig. 3. In patients with high ($n = 27$) and low ($n = 48$) sPD-L1 levels, median PFS was 5.4 months (95 % CI 4.4–7.5) and 7.2 months (95 % CI 3.7–10.3),

respectively, and high sPD-L1 correlated with shorter PFS (log-rank test, $P = 0.045$). Median OS in patients with high and low sPD-L1 levels was 12.7 months (95 % CI 9.8–15.8) and 20.4 months (95 % CI 15.8–22.9), respectively, and high sPD-L1 correlated with shorter OS (log-rank test, $P < 0.0001$). Multivariate analyses showed that there was no significant difference statistically in PFS between high and low levels of PD-L1 (HR 1.485, 95 % CI 0.6375–3.458, $P = 0.360$); meanwhile, high level of PD-L1 was significantly associated with worse OS compared with low level of PD-L1 (HR 3.307, 1.4470–7.559, $P = 0.0046$).

Discussion

Only a few reports have assessed the clinical role of serum sPD-L1 as a biomarker to predict the survival of patients with metastatic cancer (Rossille et al. 2014; Zheng et al. 2014; Cheng et al. 2015; Wang et al. 2015a, b). This study focused on evaluating the prognostic relevance of serum sPD-L1 levels in patients with metastatic GC. To the best of the authors’ knowledge, the relationship between serum sPD-L1 levels at baseline and clinical responses to first-line systemic chemotherapy in metastatic GC has not been previously investigated. As a novel finding, this study identified a negative association between OS of the first-line chemotherapy and high sPD-L1 levels at baseline. Current guidelines suggest that the eligibility criterion for antibody therapy against PD-1 or PD-L1 in metastatic GC is PD-L1 expression in tumor tissue (de Guillebon et al. 2015). Therefore, patients with high levels of sPD-L1 may obtain clinical benefit by the blockade of PD-1 or PD-L1 in metastatic GC.

The prognostic role of measuring PD-L1 expression by IHC has been reported for several tumor types, and the majority of data suggest that PD-L1 expression is

Table 2 Prognostic factors of overall survival by uni- and multivariate analyses

	Overall survival					
	Univariate analysis			Multivariate analysis		
	HR	95 % CI	P-value	HR	95 % CI	P-value
<i>Age</i>						
≤Median	1					
>Median	1.404	0.813–2.424	0.223			
<i>Gender</i>						
Female	1					
Male	0.799	0.432–1.479	0.475			
<i>ECOG PS</i>						
0–1	1					
2	1.580	0.490–5.092	0.444			
<i>Stage</i>						
Recurrence	1			1		
Stage IV	2.237	1.003–4.990	0.049	1.310	0.425–4.032	0.638
<i>Histological type</i>						
Intestinal	1					
Diffuse	1.229	0.734–2.297	0.369			
<i>HER2 status</i>						
Negative	1			1		
Positive (trastuzumab treatment)	0.537	0.293–0.983	0.044	0.880	0.438–1.767	0.720
<i>Number of metastatic sites</i>						
1	1					
≥2	1.290	0.751–2.216	0.356			
<i>Peritoneal dissemination</i>						
No	1			1		
Yes	1.699	0.974–2.963	0.062	1.397	0.769–2.540	0.272
<i>Liver metastasis</i>						
No	1					
Yes	1.046	0.608–1.800	0.870			
<i>Chemotherapeutic regimen</i>						
Dual	1					
Triplet	1.303	0.654–2.599	0.452			
<i>Serum sPD-L1 level</i>						
Low	1			1		
High	2.647	1.476–4.745	0.001	2.218	1.139–4.320	0.019

Bold values shown that *p*-value is <0.05 and there is a statistically significant difference

ECOG PS Eastern Cooperative Oncology Group Performance Status, *OS* overall survival, *CI* confidence interval, *HR* hazard ratio, *Dual* fluoropyrimidine/platinum, *Triplet* docetaxel/fluoropyrimidine/platinum, *sPD-L1* soluble programmed cell death ligand 1

associated with shorter OS (Thompson et al. 2006; Nomi et al. 2007; Kim et al. 2013; Muenst et al. 2014; Huang et al. 2015; Paydas et al. 2015; Wang et al. 2015a, b; Wu et al. 2015; Cierna et al. 2016). In gastric cancer, PD-L1 expression by IHC has been investigated by several studies, all conducted in Asian countries (Wu et al. 2006; Geng et al. 2014; Hou et al. 2014; Jiang et al. 2014; Qing et al. 2015; Kim et al. 2016). Three studies evaluated the

prognostic role of PD-L1 expression. Although interpretation of the findings is limited by differences in the definition of PD-L1 positivity and retrospective analyses, two studies indicated that high PD-L1 expression was associated with a worse prognosis in advanced GC (Wu et al. 2006; Qing et al. 2015), while the other study could not confirm the prognostic role of PD-L1 expression (Kim et al. 2016). This may be due to the heterogeneity of PD-L1 expression

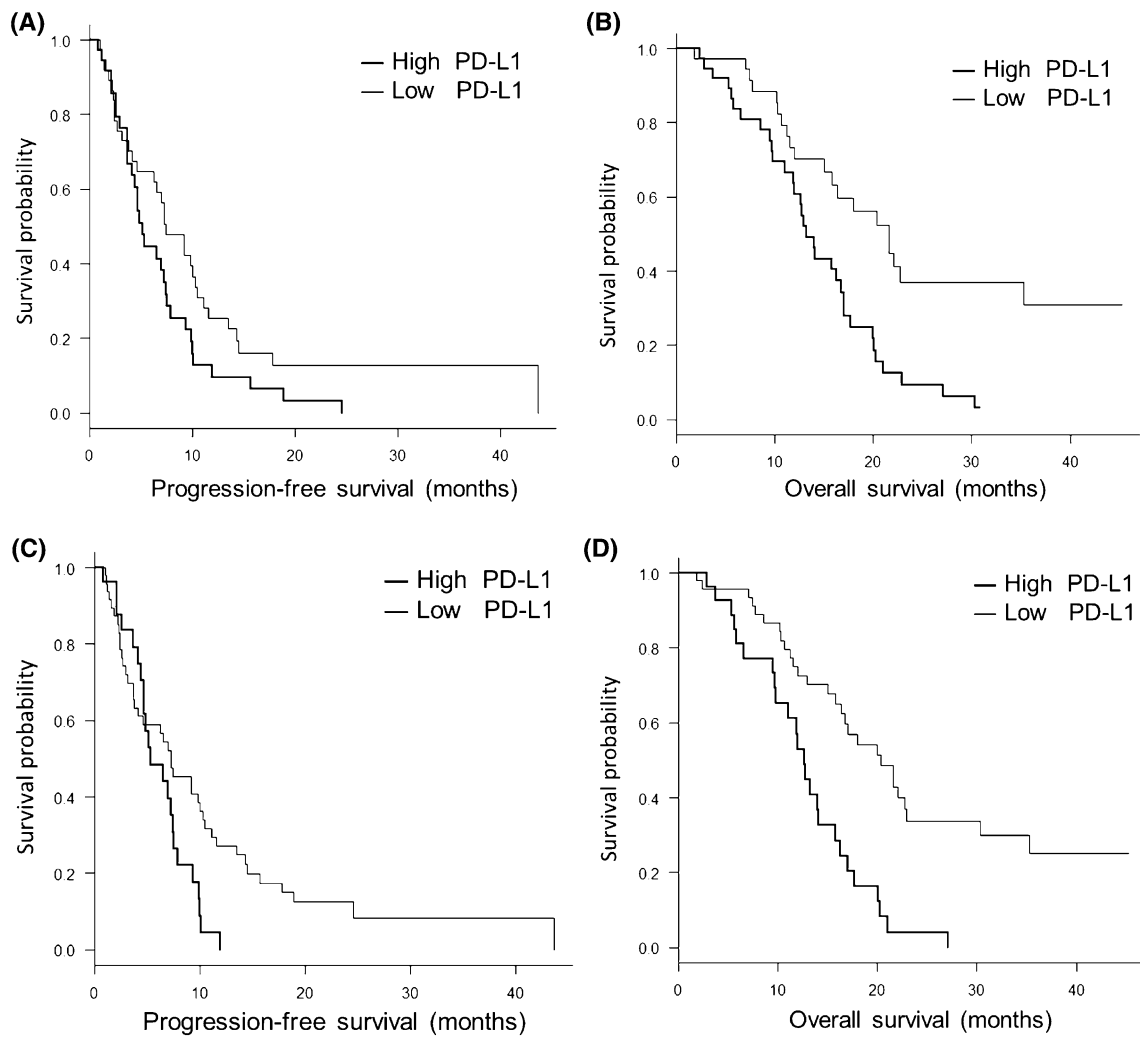


Fig. 3 Survival curves for serum sPD-L1 levels on the first-line treatment when the cutoff was the median (**a, b**) and the optimal (**c, d**) sPD-L1 value. Survival curves for progression-free survival (PFS) and overall survival (OS) are shown. High sPD-L1 was associated

with shorter OS compared with low sPD-L1 when the cutoff value was the median (**a, b**). High sPD-L1 was associated with shorter PFS and OS compared with low sPD-L1 when the cutoff value was the optimum of 1.081 ng/ml (**c, d**)

in GC, with separate tumor biopsies from the same patient producing conflicting evidence. Consistent with two of the above studies, the present study demonstrated that high sPD-L1 levels correlated with shorter OS compared with low sPD-L1 levels. This result suggests that serum sPD-L1 may be a valuable and less invasive biomarker than tumor PD-L1 expression for predicting OS.

B7-H immune checkpoint molecules usually exist as membrane-bound ligands on myeloid cells and activated lymphocytes. In previous reports, B7-H was detected in the sera of patients with malignancies such as renal cell carcinoma (Simon et al. 2006, 2007; Frigola et al. 2011). Pre-clinical data suggest that sPD-L1 is produced and released by activated mature dendritic cells and was associated with a poor prognosis in patients with renal cell carcinoma (Frigola et al. 2012). Specific binding of sPD-L1 to PD-1

has been verified by ELISA (Wang et al. 2015a, b). Taken together, these findings demonstrate that serum sPD-L1 may play an important role in the immune microenvironment of tumors. Similar to high expression of PD-L1 in tumor tissue, sPD-L1 levels detected by ELISA were significantly higher in patients with malignancies compared with healthy individuals (Rossille et al. 2014; Zheng et al. 2014; Wang et al. 2015a, b). To enable an indirect comparison of sPD-L1 levels in patients with malignancies, the present study used the same ELISA kit as that previous studies. Of note, the mean sPD-L1 level in this study (0.883 ng/ml) was very similar to that of patients with any stage of GC in another report (0.893 ng/mL) (Zheng et al. 2014). However, the predictive value of serum sPD-L1 as a prognostic marker of antibody treatment against PD-L1 or PD-1 in patients with malignancies remains to

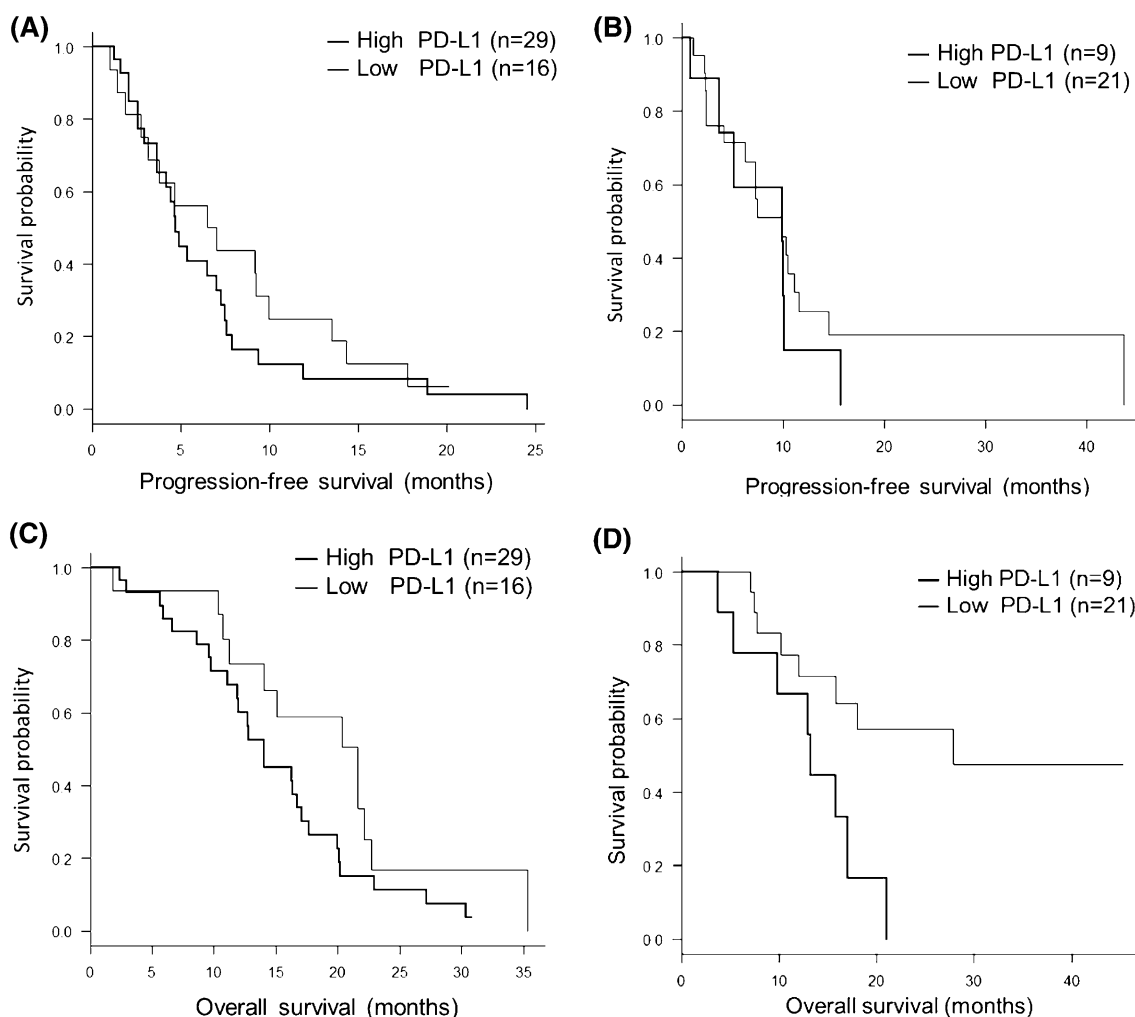


Fig. 4 Survival curves for serum sPD-L1 levels by HER2 status on the first-line treatment using median sPD-L1 as the cutoff value. Survival curves of progression-free survival (PFS) and overall survival (OS) in HER2-negative patients (**a**, **c**) and HER2-positive patients

d). OS of patients with high level of sPD-L1 tended to be associated with low level of sPD-L1 in both HER2-negative and HER2-positive patients

be clarified. With respect to anti-PD-1 antibody treatment, recent research demonstrated that serum sPD-L1 levels at pre-treatment were significantly higher in responders than in non-responders in melanoma (Dronca et al. 2015). This indicates that serum sPD-L1 may have to the potential to be not only a prognostic factor, but also a predictive marker of eligibility for anti-PD-1 antibody in metastatic GC patients.

Newly developed drugs that inhibit immune checkpoints have led to dramatic changes to the conventional treatment strategy for malignant tumors. PD-L1 is up-regulated in a variety of solid tumors, and it can be detected by IHC on both tumor and immune cells. In a phase Ib clinical trial of the anti-PD-1 antibody pembrolizumab for metastatic GC, approximately 50 % of patients achieved tumor shrinkage of target lesions and a long duration of response, despite the study population comprising patients who had

previously failed on chemotherapy (Meng et al. 2015). About 40 % of patients were diagnosed as PD-L1 positive, defined as PD-L1 staining in the stroma and ≥ 1 % of tumor cells. The definition of positivity for PD-L1 expression has not been rigorously determined in previous studies of PD-1 or PD-L1 inhibitors, and a wide spectrum of PD-L1 expression levels is observed in different tumor types (Muro et al. 2014). Reasons for the lack of clarity include differences in the cutoff value of PD-L1 positivity (proportion of positive cells: 1, 5, 10 %), positive cells (tumor cells and/or immune cells), and the manufacturer of diagnostic antibodies. Therefore, unification of IHC techniques and definitions of optimal cutoff values are required for clinical use of PD-L1 staining as a predictive marker of immune checkpoint inhibitor treatment. Recent studies have assessed immune-related biomarkers obtained by liquid biopsy (Dronca et al.

2015; Lote et al. 2015). This technique is less invasive than tissue biopsy, and it may enable the evaluation of immune status before and during treatment as opposed to archival tissue samples.

In our study, there was no difference in PFS between patients with high and low level of sPD-L1. On the other hand, patients with higher sPD-L1 level had shorter OS compared with lower sPD-L1. There were no reports of prognostic analyses of both PFS and OS according to the sPD-L1 level on the treatment of metastatic GC patients. This may indicate that sPD-L1 may be a prognostic factor but not a predictive factor of first-line treatment in metastatic gastric cancer. Combination of fluoropyrimidine and platinum is known to achieve high response rate (Bang et al. 2010; Koizumi et al. 2008, 2012; Kim et al. 2012; Yamada et al. 2015); thus, even patients with high level of sPD-L1 may obtain the clinical benefit of treatment. In addition, high WBC counts were observed in patients with high sPD-L1 level in our study. The other inflammatory findings were not associated with sPD-L1 level. To our knowledge, there were no previous reports that evaluated the association between sPD-L1 level and WBC counts of peripheral blood in patients with malignant tumors. We additionally checked leukocyte counts and neutrophil counts of peripheral blood, and neutrophil count was high in patients with high sPD-L1 level (data not shown). Neutrophil-to-leukocyte ratio (NLR) was well known as a prognostic factor in advanced gastric cancer according to previous reports (Grenader et al. 2016; Ock et al. 2016), and our study indicated that NLR was also a prognostic factors in OS by multivariate analyses in our study (data not shown). This indicated that both neutrophil counts (WBC counts) and sPD-L1 may be higher in patients with poor prognosis compared with those with better prognosis.

With respect to other biomarkers for predicting the response to anti-PD-1 or PD-L1 antibody therapy, a clinical trial in patients with GC indicated that microsatellite instability (MSI) status, a higher number of somatic mutations and antigen burden, and tumor-infiltrating lymphocytes (TILs) were considered as main candidates (Wang et al. 2013). MSI status is considered to be a particularly promising predictive biomarker of anti-PD-1 agents in gastrointestinal cancers. In colorectal cancer, pembrolizumab was associated with a high response rate in patients with MSI-high (MSI-H), whereas no responses were observed in patients with MSI-stable (MSS) (Le et al. 2015). In colorectal and non-colorectal cancers, MSI-H tumors display a higher number of somatic mutations and neo-antigen burden than MSS tumors, which strongly induces TILs and cytokine-rich tumor microenvironment, as well as expression of negative immune checkpoints, including PD-1, PD-L1, cytotoxic T-lymphocyte-associated protein 4, lymphocyte

activation gene 3, and indoleamine 2,3-dioxygenase (Le et al. 2015; Llosa et al. 2015). In advanced GC, four molecular subtypes were categorized by comprehensive evaluation of a section of the Cancer Genome Atlas project (Cancer Genome Atlas Research Network 2014). Two of the four GC subtypes, tumors positive for Epstein–Barr virus harboring *PIK3CA* mutations as well as amplification of *JAK2*, *CD274* and *PDCD1LG2* (9 %) and microsatellite unstable tumors with elevated gene mutation rates (22 %), were considered to be vulnerable to blockade of PD-1 or PD-L1.

The present study has several limitations. The potential for selection bias cannot be excluded, and this was an exploratory study with a limited sample size. Patients with insufficient serum samples for ELISA were excluded, while a higher proportion of HER2-positive than HER2-negative patients were included. Of note, age and ECOG PS, which are well-known prognostic factors, were not included in multivariate analysis because these covariates had P -values ≥ 0.2 in univariate analysis. Furthermore, prospective data are required to validate the results of this study. Considering the heterogeneity of PD-L1 expression in GC tissue, it is necessary to determine the level of concordance between PD-L1 expression by IHC and serum sPD-L1 by ELISA.

This study revealed that the detection of high levels of serum sPD-L1 was associated with a poor survival prognosis in patients with metastatic GC. Although the potential for differences between sPD-L1 in blood and PD-L1 expression in tissue remains unclear, it is possible that serum sPD-L1 is a useful and less invasive biomarker in metastatic GC patients. Serum level of sPD-L1 in 40 % of enrolled patients was lower than measurable range in our study, and more sensitive ELISA kit may be required for strict quantitative analyses. Further investigations are warranted to confirm the relevance of sPD-L1 as a prognostic or predictive biomarker of systemic chemotherapy as well as immune checkpoint inhibitors.

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Compliance with ethical standards

Conflict of interest The authors declare that they have no conflict of interest.

Ethical approval All procedures performed were in accordance with the ethical standards of the national regulations and with the 1964 Declaration of Helsinki and its later amendments or comparable ethical standards.

Informed consent Informed consent of Biobank of National Cancer Center and using the clinical data was obtained from all individual participants included in the study.

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