



Tumor PD-L1 expression is associated with outcomes in stage III non-small cell lung cancer (NSCLC) patients treated with consolidation durvalumab

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Background: Durvalumab is an anti-PD-L1 immune checkpoint inhibitor approved for consolidation therapy for patients with stage III non-small cell lung cancer (NSCLC) after chemoradiation. The purpose of our study was to evaluate the association between the degree of tumor PD-L1 expression and outcomes of stage III NSCLC patients treated with durvalumab.

Methods: We conducted a retrospective analysis of all the patients who received durvalumab between July 2017 and July 2019 at our facilities and were diagnosed with or progressed to stage III NSCLC before durvalumab consolidation. Patients were divided into groups based on the degree of PD-L1 expression: <1%, 1–49%, and 50–100%. Overall survival and progression-free survival were estimated by the Kaplan-Meier method and the Multivariate Cox proportional hazard model was used to assess the effect of PD-L1 expression level on OS and PFS, adjusting for age and gender.

Results: We identified 121 patients with stage III NSCLC that underwent durvalumab consolidation. Of them, 29.8% had PD-L1 expression of 50–100%, 24.8% had PD-L1 expression of 1–49%, and 27.3% had PD-L1 expression of <1%, while 18.2% were not tested for PD-L1 expression. The rate of cancer progression in the group with 50–100% PD-L1 expression was 16.7% compared to 60% in the 1–49% expression group and 54.6% in the <1% expression group, and the 1-year survival rates were higher in the 50–100% group (97%) compared to the 1–49% group and the <1% group (73% and 78%, respectively; $P=0.028$). Survival analysis via Kaplan-Meier revealed a significant difference in both PFS ($P<0.0001$) and OS ($P<0.028$) based on the extent of PD-L1 expression. Multivariate analysis revealed that a PD-L1 expression >50% was the only factor that was significantly associated with improved PFS (HR =0.205, $P=0.0004$) and OS (HR =0.339, $P=0.04$).

Conclusions: Our study demonstrated that patients whose tumors had >50% PD-L1 expression had significantly longer progression-free survival and overall survival than those with lower PD-L1 expression. This suggests that the degree of tumor PD-L1 expression may play a role in predicting benefit from durvalumab for these patients.

Keywords: Lung cancer; durvalumab; programmed-death ligand 1 (PD-L1)

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Introduction

Lung cancer is estimated to be the number one cause of cancer-related deaths in the US in 2021, and non-small cell lung cancer (NSCLC) accounts for about 85% of lung cancer cases (1). Given the findings of the PACIFIC trial in which durvalumab improved both the progression-free survival (PFS) and overall survival (OS) of patients with stage III NSCLC who had completed chemotherapy and radiotherapy, durvalumab consolidation has now become a mainstay of treatment for patients with stage III NSCLC (2-4). Durvalumab is a monoclonal antibody that targets the programmed-death ligand 1 (PD-L1) on tumor cells, allowing them to be recognized and killed by the immune system (5,6). Testing for PD-L1 expression is typically done via immunohistochemistry (IHC) assays; some of the more common monoclonal antibody clones in testing kits include the 22C3, SP263, and the 28-8 clones (7).

There is little research to indicate the role of PD-L1 expression in predicting outcomes of patients being treated with durvalumab. Several studies indicate that higher PD-L1 expression confers better treatment outcomes in NSCLC patients treated with pembrolizumab (8,9) atezolizumab (10,11) and nivolumab (12), yet other studies have suggested otherwise (13). As for durvalumab, there is data to suggest that PD-L1 expression is associated with higher response rates (14,15). However, the issue with most of these studies is that they are phase I/II with small sample sizes. Additionally, there are different cutoffs used to determine “high” PD-L1 expression. In the PACIFIC trial, patients were divided into groups of those with <25% of tumor cells expressing PD-L1 and those with \geq 25% PD-L1 expression. Improvement in PFS and OS were seen with durvalumab regardless of the extent of PD-L1 expression, although it seemed that the improvement in OS was least noted in patients with <1% PD-L1 expression (2-4). The purpose of our study was to evaluate the association between the degree of PD-L1 expression and outcomes of stage III NSCLC patients treated with durvalumab. We present the following article in accordance with the STROBE reporting checklist (16) (available at <https://dx.doi.org/10.21037/tlcr-21-249>).

Methods

This was a retrospective study of patients with stage III NSCLC treated with durvalumab within the Cleveland Clinic Foundation enterprise. We used our institutional

pharmacy records to identify all the patients who received durvalumab between July 2017 and July 2019 at our facilities and selected those who were diagnosed with or progressed to Stage III NSCLC before durvalumab consolidation. Patients who had small cell lung cancer, were treated at stage IV of their disease, or who only received a portion of their care here for a second opinion and then were lost to follow up as they continued their care at other institutions were excluded from our analyses. We decided to select all of the stage III NSCLC patients treated with durvalumab to minimize selection bias and have the most data possible. Information was collected from the patients' electronic medical charts and stored and managed using a secure database between July 2019 and September 2020.

The study was conducted in accordance with the Declaration of Helsinki (as revised in 2013). The study was approved by the institutional review board of the Cleveland Clinic Foundation (NO.: 14-562) and individual consent for this retrospective analysis was waived. Information that was collected about each patient included demographics (age, race, sex), smoking history, clinical and pathological information regarding their cancer (histological subtype, initial stage at diagnosis, T/N/M status, tumor marker expression, PD-L1 expression), treatment course (number of durvalumab doses, side effects), and clinical outcomes.

Statistical analysis

PFS was defined as the time from initiation of durvalumab until cancer progression (per RECIST criteria) or recurrence was identified (or until the last day durvalumab was known to be effective). OS was defined as the time from initiation of durvalumab until the last day the patient was known to be alive.

Patients were divided into groups based on the degree of PD-L1 expression: <1% (PD-L1 absent), 1-49% (low expression), and 50-100% (high expression). The Cleveland Clinic Laboratory performs PD-L1 testing on tissue samples via immunohistochemistry testing for the Dako 22C3 PD-L1 clone to assess the proportion of tumor cells that express PD-L1. Patient characteristics were summarized in median and range for continuous variables, and in frequencies and percentages for categorical variables. Fisher's exact test was used to associate patient characteristics with PD-L1 expression status. Wilcoxon rank-sum test was used to compare age, pack-year, and the number of doses between PD-L1 groups. Overall survival and progression-free survival were estimated by the Kaplan-

Meier method and were compared using log-rank testing between patient groups. Multivariate Cox proportional hazard model was used to assess the effect of PD-L1 expression level on OS and PFS, adjusting for age and gender. All tests were two-sided and P values of 0.05 or less were considered statistically significant. Statistical analysis was carried out using SAS Studio 3.7 (SAS Institute, Cary, NC) and R 3.6 (R Foundation, Vienna, Austria).

Results

We identified 138 patients treated with durvalumab at our facilities, of which 121 had stage III NSCLC and fit our

forementioned research criteria. 50.4% were female, and the mean age was 68.3 years. Of them, 29.8% had PD-L1 expression of 50–100%, 24.8% had PD-L1 expression of 1–49%, and 27.3% had PD-L1 expression of <1%, while 18.2% were not tested for PD-L1 expression (*Table 1*). There was no significant difference in age, sex, smoking pack-years, the histologic subtype of NSCLC, and the number of durvalumab doses administered between patients who expressed PD-L1 and those who did not.

Cancer progression occurred in 16.7% of the patients in the group with 50–100% PD-L1 expression compared to 60% in the 1–49% expression group and 54.6% in the <1% expression group, and the 1-year survival rates were higher

Table 1 Summary of patient characteristics by PD-L1 status

Patient characteristics	PD-L1 extent						All		P value
	<1%		1–49%		50–100%		N	%	
	N	%	N	%	N	%			
Sex									
Female	21	63.64	12	40	16	44.44	49	49.49	0.13
Male	12	36.36	18	60	20	55.56	50	50.51	
Race									
Others	7	21.21	6	20	9	25	22	22.22	0.91
White	26	78.79	24	80	27	75	77	77.78	
Smoking status									
Unknown	0	0	1	3.33	0	0	1	1.01	0.12
Current smoker	14	42.42	5	16.67	8	22.22	27	27.27	
Former smoker	17	51.52	23	76.67	27	75	67	67.68	
Never smoker	2	6.06	1	3.33	1	2.78	4	4.04	
Tumor histology									
Adenocarcinoma	14	42.42	19	63.33	18	50	51	51.52	0.13
Squamous cell carcinoma	17	51.52	9	30	15	41.67	41	41.41	
Large cell carcinoma	0	0	2	6.67	0	0	2	2.02	
Other	2	6.06	0	0	3	8.33	5	5.05	
Clinical stage at diagnosis									
I/II	4	12.12	3	10	5	13.89	12	12.12	0.41
IIIA	12	36.36	9	30	17	47.22	38	38.38	
IIIB	16	48.48	16	53.33	10	27.78	42	42.42	
IIIC	1	3.03	2	6.67	4	11.11	7	7.07	

Table 1 (continued)

Table 1 (continued)

Patient characteristics	PD-L1 extent						All		P value
	<1%		1–49%		50–100%		N	%	
	N	%	N	%	N	%			
TNM, T status at diagnosis									
Unknown	0	0	1	3.33	0	0	1	1.01	0.16
T1	6	18.18	13	43.33	12	33.33	31	31.31	
T2	11	33.33	5	16.67	9	25	25	25.25	
T3	9	27.27	10	33.33	9	25	28	28.28	
T4	7	21.21	1	3.33	5	13.89	13	13.13	
Tx	0	0	0	0	1	2.78	1	1.01	
TNM, N status at diagnosis									
Unknown	0	0	1	3.33	0	0	1	1.01	0.11
N0	6	18.18	0	0	5	13.89	11	11.11	
N1	5	15.15	7	23.33	6	16.67	18	18.18	
N2	15	45.45	12	40	20	55.56	47	47.47	
N3	7	21.21	10	33.33	5	13.89	22	22.22	
TNM, M status at diagnosis									
Unknown	0	0	1	3.33	0	0	1	1.01	98.99
M0	33	100	29	96.67	36	100	98		
EGFR mutation status									
Unknown	19	57.58	7	23.33	15	41.67	41	41.41	2.02
Absent	13	39.39	22	73.33	21	58.33	56	56.57	
Present	1	3.03	1	3.33	0	0	2		
ALK mutation status									
Unknown	21	63.64	12	40	17	47.22	50	50.51	49.49
Absent	12	36.36	18	60	19	52.78	49		
ROS mutation status									
Unknown	22	66.67	15	50	21	58.33	58	58.59	41.41
Absent	11	33.33	15	50	15	41.67	41		
KRAS mutation status									
Unknown	23	69.7	12	40	18	50	53	53.54	12.12
Absent	9	27.27	15	50	10	27.78	34	34.34	
Present	1	3.03	3	10	8	22.22	12		
BRAF mutation status									
Unknown	22	66.67	12	40	17	47.22	51	51.52	2.02
Absent	11	33.33	16	53.33	19	52.78	46	46.46	
Present	0	0	2	6.67	0	0	2		

Table 1 (continued)

Table 1 (continued)

Patient characteristics	PD-L1 extent						All		P value
	<1%		1–49%		50–100%		N	%	
	N	%	N	%	N	%			
MET mutation status									
Unknown	23	69.7	14	46.67	17	47.22	54	54.55	
Absent	10	30.3	16	53.33	19	52.78	45	45.45	
RET mutation status									
Unknown	24	72.73	18	60	24	66.67	66	66.67	
Absent	8	24.24	12	40	12	33.33	32	32.32	
Present	1	3.03	0	0	0	0	1	1.01	
BRCA1 mutation status									
Unknown	32	96.97	29	96.67	35	97.22	96	96.97	
Absent	1	3.03	1	3.33	1	2.78	3	3.03	
ERBB2 mutation status									
Unknown	24	72.73	19	63.33	25	69.44	68	68.69	
Absent	9	27.27	11	36.67	11	30.56	31	31.31	
Cancer progression									
Yes	18	54.55	18	60	6	16.67	42	42.42	
No	15	45.45	12	40	30	83.33	57	57.58	
Treatment discontinued due to progression									
Yes	10	30.3	12	40	5	13.89	27	27.27	0.048
No	23	69.7	18	60	31	86.11	72	72.73	
Treatment discontinued due to adverse events									
Yes	11	33.33	7	23.33	8	22.22	26	26.26	0.55
No	22	66.67	23	76.67	28	77.78	73	73.74	
Treatment discontinued due to other reasons									
Yes	2	6.06	4	13.33	3	8.33	9	9.09	0.62
No	31	93.94	26	86.67	33	91.67	90	90.91	
Living status									
Living	23	69.7	18	60	31	86.11	72	72.73	
Deceased	10	30.3	12	40	5	13.89	27	27.27	
All	33	100	30	100	36	100	99	100	

P values by Fisher's exact test. Patients with unknown PD-L1 status were not included in this table. PD-L1, programmed-death ligand 1; TNM, tumor node metastases; EGFR, epidermal growth factor receptor oncogene; ALK, anaplastic lymphoma kinase oncogene; ROS, Ros oncogene; KRAS, KRAS oncogene; BRAF, BRAF oncogene; MET, MET oncogene; RET, RET oncogene; BRCA, breast cancer oncogene; ERBB2, ERBB2 oncogene.

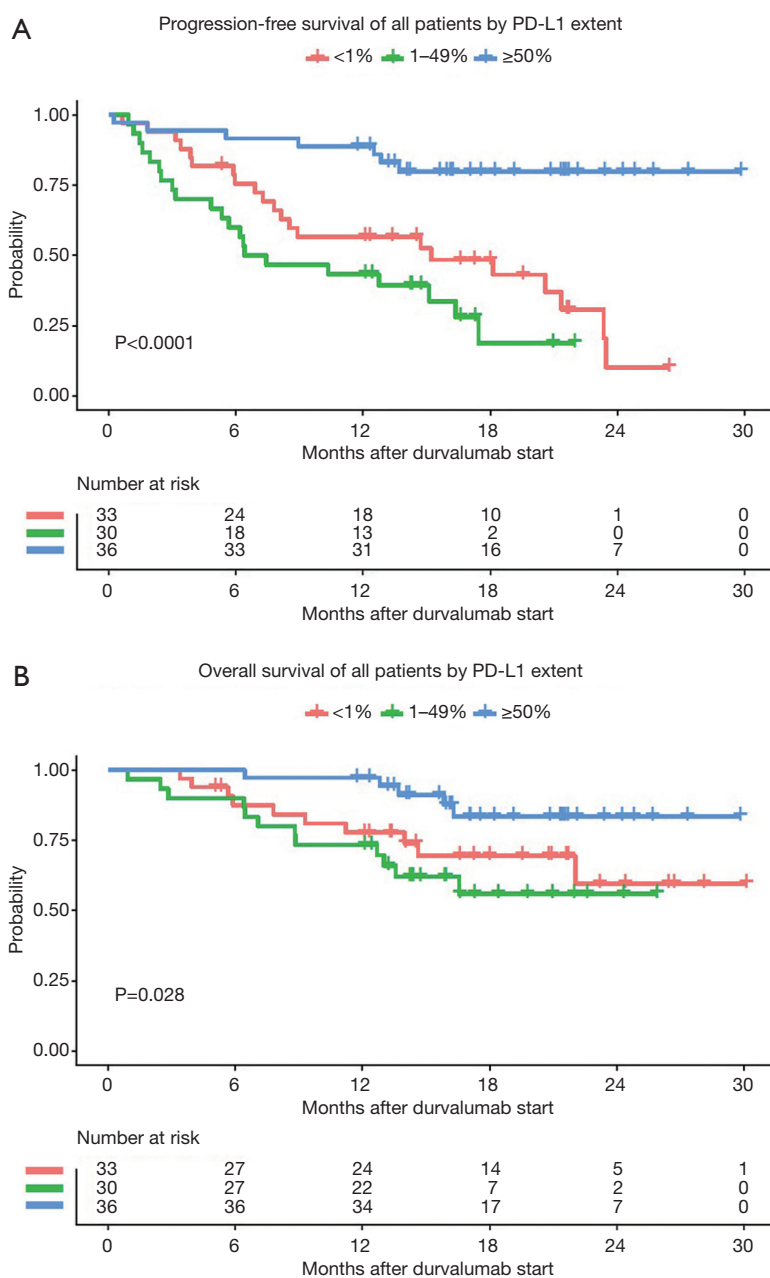


Figure 1 Kaplan-Meier curves of (A) progression-free survival and (B) overall survival by the extent of tumor PD-L1 expression. Survival was compared using log-rank testing between patient groups. PD-L1, programmed-death ligand 1.

in the 50–100% group (97%) compared to the 1–49% group and the <1% group (73% and 78%, respectively; $P=0.028$).

The median PFS for the group with 50–100% expression was 16.9 months, compared to 7.0 months for 1–49% expression group and 12.5 months for the <1% expression group. The median OS was 17.6 months for the 50–100%

expression group, 14.5 months for the 1–49% expression group, and 14.8 months for the <1% expression group. Survival analysis via Kaplan-Meier revealed a significant difference in both PFS (log-rank $P < 0.0001$) and OS (log-rank $P < 0.028$) based on the extent of PD-L1 expression (Figure 1). Multivariate analysis revealed that a PD-L1 expression >50% was the only factor that was significantly

Table 2 Summary of multivariate Cox proportional hazard model for PFS and OS

	Factor	Comparison	Hazard ratio	95% LCL	95% UCL	P value
PFS	PD-L1	1–49% vs. <1%	1.446	0.752	2.777	0.2686
		50–100% vs. <1%	0.205	0.086	0.491	0.0004
		50–100% vs. 1–49%	7.037	2.896	17.096	<0.0001
OS	PD-L1	1–49% vs. <1%	1.289	0.535	3.176	0.5719
		50–100% vs. <1%	0.339	0.104	0.973	0.04
		50–100% vs. 1–49%	3.807	1.336	10.849	0.01

Age and gender were forced into model to adjust their effects on PFS and OS. PD-L1 was significant after adjusting for them. PD-L1, programmed-death ligand 1; PFS, progression-free survival; OS, overall survival; LCL, lower control limit; UCL, upper control limit.

associated with improved PFS (HR =0.205, P=0.0004) and OS (HR =0.339, P=0.04) when evaluated with age, sex, race, smoking status, the histologic subtype of NSCLC, tumor size and lymph node status (Table 2).

Conclusions

The results of our study demonstrate that the degree of tumor PD-L1 expression may predict the response of NSCLC patients to durvalumab therapy and that patients with >50% PD-L1 expression had better outcomes and survived longer than those with <50% expression. Notably, patients with PD-L1 expression of 1–49% and <1% had relatively similar outcomes, which may explain why studies that used the cutoff of <25% and ≥25% PD-L1 expression did not identify as significant a difference in outcomes (2). Limitations of our study include its retrospective design with potential resulting bias and our sample size of 121 patients, and external validity of our findings would require regular PD-L1 testing in the workup of NSCLC. Yet to our knowledge, this the largest observational study that showed a clear survival benefit favoring higher PD-L1 expression for stage III NSCLC patients undergoing durvalumab consolidation. These findings could potentially affect clinical decision-making regarding therapy selection and monitoring of these patients. More research is needed to determine the relationship between the extent of PD-L1 expression and the treatment outcomes in patients with locally advanced NSCLC. We recommend that future trials revolving around the use of immunotherapy, particularly durvalumab, in patients with NSCLC pay attention to the extent of PD-L1 expression and use the above cutoffs to better categorize patients and study their outcomes.

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Footnote

Reporting Checklist: The authors have completed the STROBE reporting checklist. Available at <https://dx.doi.org/10.21037/tlcr-21-249>

Data Sharing Statement: Available at <https://dx.doi.org/10.21037/tlcr-21-249>

Conflicts of Interest: All authors have completed the ICMJE uniform disclosure form (available at <https://dx.doi.org/10.21037/tlcr-21-249>). NAP has served as a consultant on advisory boards for Astrazeneca, Merck, BMS, Genentech, and Eli Lilly, with payments made to him. The other authors have no conflicts of interest to disclose.

Ethical Statement: The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved. The study was conducted in accordance with the Declaration of Helsinki (as revised in 2013). The study was approved by the institutional review board of the Cleveland Clinic Foundation (NO: 14-562) and individual consent for this retrospective analysis was waived.

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