ORIGINAL RESEARCH

Altered status of programmed death-ligand 1 after recurrence in resected lung adenocarcinoma patients

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Purpose: Programmed death-ligand 1 (PD-L1) is found to be overexpressed in non-small cell lung cancer. The present study intended to evaluate the status of PD-L1 expression in patients with resection and recurrent lung adenocarcinoma.

Patients and methods: Matched resection and recurrent tumor samples were harvested from 65 lung adenocarcinoma patients. Immunohistochemistry was used to evaluate the status of PD-L1 expression. Kaplan–Meier method was used for survival analysis.

Results: A total of 65 patients of lung adenocarcinoma were enrolled. They underwent complete resection and had recurrence after adjuvant treatment. PD-L1 expression was identified in 43.1% (28/65) of resection samples vs 55.4% (36/65) of recurrent samples. Ten patients shifted from negative to positive, whereas another two samples showed the opposite. Patients with PD-L1 expression showed worse disease-free survival than the PD-L1-negative counterparts. The expression of PD-L1 in recurrent samples was a significant favorable factor for epidermal growth factor receptor tyrosine kinase inhibitors (*EGFR*-TKIs) (11.2 vs 8.2 months, *P*=0.030).

Conclusion: The status of PD-L1 expression may alter between resection and recurrent samples. Also, the status of PD-L1 expression after recurrence is a better prognostic factor for *EGFR*-TKIs.

Keywords: programmed death-ligand 1, overexpression, lung adenocarcinoma, survival

Introduction

Non-small cell lung cancer accounts for >80% of all lung carcinomas and lung adenocarcinoma is predominant.[1] At present, chemotherapy, radiotherapy, molecular therapy and surgery are the major treatment modalities for lung cancer treatment.[2] However, most patients remain resistant to treatment with a median survival time of <2 years.[3–7]

It is well known that immune therapy based on programmed death-1 (PD-1) and its ligand PD-L1 shows promising efficacy.[8,9] The progression-free survival (PFS) was longer in patients on immunotherapy than those on chemotherapy based on high-expression PD-L1.[10] PD-L1 has been identified as a useful biomarker for immunotherapy. Chemotherapy could alter the status of PD-L1 expression in squamous cell carcinoma.[11] However, PD-L1 status is not thoroughly examined for lung adenocarcinoma in resection and recurrent samples.

The present study was intended to detect the PD-L1 expression status in resection and recurrent samples and validate its clinical value in lung adenocarcinoma.

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Materials and methods

Patient selection

During the period 2009–2015, 65 patients with pathologically confirmed lung adenocarcinoma were recruited from Yinzhou Hospital of Zhejiang Province. The inclusion criteria were as follows: 1) pathologically proven primary lung adenocarcinoma with resection and recurrent samples and 2) no prior local treatment in recurrent samples. Histologic typing was confirmed as lung adenocarcinoma according to the 2004 World Health Organization classification scheme. Also, TNM stage was determined according to the 2009 criteria. The Ethics Committee of Yinzhou Hospital approved this study (IRB-2015-027), and written informed consent was obtained from each participant.

Immunohistochemical analysis of PD-L1 expression

Analysis of PD-L1 expression was performed on 4–6 μm thick, formalin-fixated, paraffin-embedded tissue with immunohistochemical staining. PD-L1 from Proteintech Group Inc. (Chicago, IL, USA) was utilized. UltraVision Quanto Detection System (Thermo Fisher Scientific, Waltham, MA, USA) was used for detecting PD-L1 expression according to the manufacturer's instructions.

Semi-quantitative H-score (a maximal value of 300 corresponding to 100% of tumor cells staining positive for PD-L1 with an overall staining intensity score of 3) was determined by multiplying the percentage of stained cells by an intensity score (0, absent; 1, weak; 2, moderate; 3, strong). A 5% proportion of membranous staining of tumor cells with H-score ≥5 was designated as the cutoff for PD-L1 positivity. Three independent pathologists assessed the expression of PD-L1 status.

Efficacy evaluation

Tumor efficacy was evaluated according to the Response Evaluation Criteria in Solid Tumors (RECIST 1.1). Tumor responses included complete response, partial response, stable disease and progressive disease. Also, disease control rate was defined as a combination of objective responses and stabilization. The median follow-up period was 38 (13–71) months. The last follow-up date was October 31, 2016.

Statistical analyses

Chi-squared test was used for evaluating the relationship between clinical characteristics and status of PD-L1 expression. Survival curves were calculated by the Kaplan–Meier method. Statistical analysis was performed with SPSS 16 software (SPSS Inc., Chicago, IL, USA). $P < 0.05$ was deemed as statistically significant.

Results

Clinical characteristics

A total of 65 patients of lung adenocarcinoma were recruited. There were 37 males and 28 females with a median age of 62 (35–75) years. The smoking status was former/current (n=34) and never (n=31) smokers. All patients underwent complete resection. Their pathologic stages were I (n=19), II (n=15) and IIIA (n=32). Among the patients, 56 patients received adjuvant treatment, including chemotherapy (n=25), radiotherapy (n=3) and chemoradiotherapy (n=28). Their clinicopathologic characteristics are summarized in Table 1.

PD-L1 status in resection and recurrent samples

As for PD-L1 expression, 28 (43.1%) patients were positive in resection samples vs 36 in recurrent samples. Ten patients shifted from negative to positive, whereas another two samples showed the opposite (P=0.005; Table 2).

No correlations existed in gender, age, smoking status or status of PD-L1 expression. Similarly, there was no correlation between clinicopathologic characteristics and PD-L1 expression in recurrent samples. The correlations

Table 1 Clinical characteristics of the patient population (N=65)

Variable	Number
Gender	
Male	37
Female	28
Age, years	
Median	62
<65	47
≥65	18
Smoking status	
Never	31
Former/current	34
Pathological stage after surgery	
I + II	33
IIIA	32
EGFR status	
Mutation	28
Wild type	24
Unknown	13
PD-L1 expression in resected samples	
Positive	28
Negative	37
PD-L1 expression in recurrence samples	
Positive	36
Negative	29

Abbreviations: *EGFR*, epidermal growth factor receptor; PD-L1, programmed death-ligand 1.

Table 2 Clinical characteristics of patients with altered PD-L1 expression

Case	Gender/ age (years)	Pathological stage	Smoking status	PD-L1 expression		Disease-free survival, months	Overall survival, months
				Resected samples	Recurrence samples		
1	M/61	IIIA	Former	Positive	Negative	11.5	37.9
2	M/55	IIIA	Never	Positive	Negative	20.0	45.5
3	M/67	IIIA	Former	Negative	Positive	26.0	55.0
4	M/59	IIB	Former	Negative	Positive	19.0	38.0
5	M/52	IIIA	Current	Negative	Positive	7.0	21.0
6	M/71	IB	Former	Negative	Positive	39.0	57.2
7	F/60	IIIA	Never	Negative	Positive	12.5	43.0
8	M/63	IIA	Never	Negative	Positive	33.0	49.0+
9	F/66	IB	Never	Negative	Positive	41.5	61.6+
10	M/51	IIIA	Current	Negative	Positive	16.0	36.5
11	F/38	IIB	Current	Negative	Positive	13.3	35.2
12	F/51	IIIA	Never	Negative	Positive	14.5	55.4

Abbreviation: PD-L1, programmed death-ligand 1.

PD-L1 expression and gene mutations

Among the patients, the expression of epidermal growth factor receptor (*EGFR*) and *ALK* was detected in 52 patients. There were 28 *EGFR*-mutant and 24 wild-type patients. Four harbored *ALK* rearrangement and 48 were *ALK* negative. The results of gene mutations were based on resection samples. The frequencies of PD-L1 expression were 66.7% and 42.9% in *EGFR*-mutant and wild-type patients, respectively (*P*=0.08). No correlations existed between PD-L1 expression and *ALK* rearrangements (*P*=0.72).

Treatment

All the patients had a postoperative onset of recurrence or metastasis. The median values of disease-free survival and overall survival were 22.0 months (95% confidence interval: 19.1–26.2) and 37.1 months (95% confidence interval: 34.0–39.4), respectively. All recurrences were treated. All 28 *EGFR*-mutant patients received *EGFR*-TKIs. According to the status of PD-L1 expression in recurrent samples, a longer PFS was observed in PD-L1-positive than PD-L1-negative patients (11.2 vs 8.2 months, *P*=0.030; Figure 1). Also, 55 patients received first-line chemotherapy. No correlations of efficacy difference existed between PD-L1-positive and PD-L1-negative patients (4.3 vs 4.6 months, *P*=0.57).

Survival analyses

Based upon PD-L1 expression in resection samples, a trend of longer disease-free survival existed in PD-L1-negative than PD-L1-positive patients (23.0 vs 18.8 months, *P*=0.01; Figure 2). However, there was no significant difference in

Table 3 Correlation between PD-L1 expression and clinical characteristics

Variable	Resected samples			Recurrence samples		
	PD-L1 positive	PD-L1 negative	*P*-value	PD-L1 positive	PD-L1 negative	*P*-value
Gender			0.64			0.45
Male	15	22		19	18	
Female	13	15		17	11	
Age (years)			0.21			0.049
<65	18	29		22	25	
≥65	10	8		14	4	
Smoking status			0.49			0.56
Never	12	19		16	15	
Former/current	16	18		20	14	
Pathologic stage			0.16			0.44
I + II	17	16		NA	NA	NA
IIIA	11	21		NA	NA	NA
EGFR mutation			0.086			
Yes	16	12		NA	NA	NA
No	8	16		NA	NA	NA

Abbreviations: *EGFR*, epidermal growth factor receptor; NA, not applicable; PD-L1, programmed death-ligand 1.
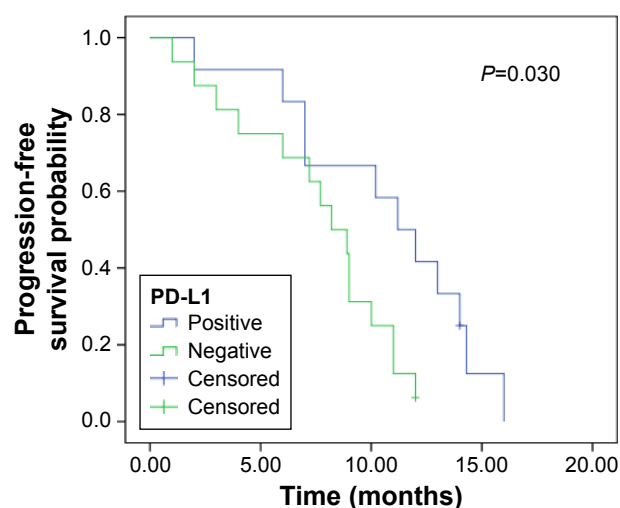
Figure 1 Progression-free survival of *EGFR*-TKIs in *EGFR*-mutant patients according to the status of PD-L1 expression in recurrent samples.
Abbreviations: *EGFR*, epidermal growth factor receptor; PD-L1, programmed death-ligand 1; TKIs, tyrosine kinase inhibitors.

overall survival according to the status of PD-L1 expression in resection samples (39.0 vs 34.0 months, *P*=0.225).

Similarly, no survival difference existed in recurrent samples between PD-L1-positive and PD-L1-negative groups (39.0 vs 34.5 months, *P*=0.226).

Discussion

The status of PD-L1 expression might alter after recurrence in some completely resected lung adenocarcinoma patients. The expression of PD-L1 became upregulated in recurrent samples. The status of PD-L1 after recurrence could better

Figure 2 Comparison of disease-free survival based upon the status of PD-L1 expression in resection lung adenocarcinoma samples.
Abbreviation: PD-L1, programmed death-ligand 1.

predict the favorable efficacy of *EGFR*-TKIs. To the best of our knowledge, this was the first study of detecting differences between resection and recurrent samples in lung adenocarcinoma patients.

Many solid carcinomas were found to show overexpression of PD-L1.[12–16] It is well known that gene status might be affected by chemotherapy.[17] However, the status of PD-L1 could be altered over time. PD-L1 status might alter before and after dosing of *EGFR*-TKIs in lung adenocarcinoma patients.[18] Also, the magnitude of PD-L1 expression increased after chemotherapy in lung squamous cell carcinoma patients.[11] In the present study, the frequency of PD-L1 expression increased in several recurrent samples. One reason for this might be an alteration of immune microenvironment after recurrence. Also, it was due to heterogeneity of tumors between resection and recurrent samples. Last but not least, most patients received adjuvant chemotherapy or radiotherapy, thus affecting PD-L1 expression partially. Because of inconstancy of PD-L1 expression in resection and recurrent samples, the status of PD-L1 expression should be remonitored in patients with recurrent or metastatic samples.

PD-L1 serves as a favorable biomarker for the efficacy of *EGFR*-TKIs,[19] since *EGFR*-TKIs can inhibit tumor cell viability and indirectly enhance antitumor immunity through downregulation of PD-L1.[20] In the present study, a longer PFS existed in PD-L1-positive patients than the negative counterparts. As concluded from previous and present studies, PD-L1 expression might affect the clinical efficacy of *EGFR*-TKIs. Prospective studies with a larger number of patients should be performed.

Limitations

The present study had some inherent limitations. First, it was retrospective in nature, and the sample size was too small. Second, use of only one antibody might have influenced the frequency of PD-L1 expression. Third, only 28 *EGFR*-mutant patients received *EGFR*-TKIs, so few cases were available for efficacy analysis. Thus, our results should be validated by future studies of a larger sample size. However, our results may carry clinical implications for immunotherapy in lung adenocarcinoma patients.

Conclusion

PD-L1 expression may be upregulated after recurrence in resection lung adenocarcinoma patients. Also, PD-L1 may be a useful biomarker for predicting the efficacy of *EGFR*-TKIs.
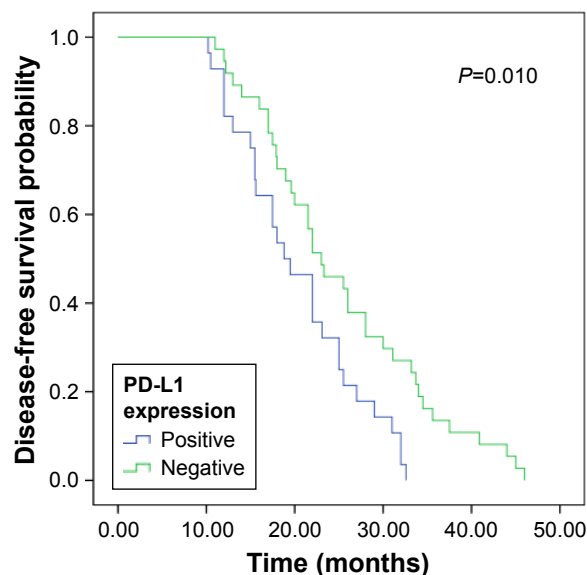
Acknowledgment

Disclosure

The authors report no conflicts of interest in this work.

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