

# Pretreatment Lung Immune Prognostic Index Is a Prognostic Marker of Chemotherapy and Epidermal Growth Factor Receptor Tyrosine Kinase Inhibitor

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## Abstract

**Background:** Lung immune prognostic index (LIPI) was recently developed on the basis of the combination of baseline derived neutrophil to lymphocyte ratio (dNLR) and lactate dehydrogenase (LDH). This index was demonstrated as a specific biomarker of immune checkpoint inhibitors for non-small cell lung cancer (NSCLC). We aimed to show that LIPI may be a useful biomarker of cytotoxic chemotherapy and epidermal growth factor receptor (EGFR) tyrosine kinase inhibitor (TKI) for NSCLC.

**Methods:** We retrospectively collected 175 wild-type EGFR adenocarcinomas, 131 NSCLCs harboring mutant EGFR and 110 squamous cell carcinomas. All patients initiated first-line cytotoxic chemotherapy or EGFR-TKI monotherapy between July 2007 and August 2017 at our hospital. These patients were divided into good, intermediate and poor LIPI groups. We compared their overall survival (OS) and progression-free survival (PFS). Multivariate analyses detected prognostic and predictive factors of OS and PFS.

**Results:** The good LIPI group survived longer than the intermediate and poor LIPI groups in wild-type EGFR adenocarcinoma (good, intermediate and poor LIPI groups: median 19.6, 11.5 and 3.3 months,  $P < 0.01$ , respectively) and mutant EGFR NSCLC (45.4, 25.6 and 15.7 months,  $P < 0.01$ ). The PFS of good LIPI group was significantly longer than those of the other two groups in mutant EGFR NSCLC (16.6, 12.6 and 8.3 months,  $P < 0.01$ ). The intermediate group (hazard ratio (HR) 1.49, 95% confidential interval (CI) 1.03 - 2.15,  $P = 0.04$ ) of wild-type EGFR adenocarcinoma, intermediate (HR 2.30, 95% CI 1.33 - 3.99,  $P < 0.01$ ) and poor (HR 2.76, 95% CI 1.03 - 7.42,  $P = 0.04$ ) groups of mutant EGFR NSCLC were independent prognostic factors of poor OS. The intermediate (HR 1.57, 95% CI 1.01 - 2.44,  $P = 0.04$ ) and poor (HR 2.63, 95% CI 1.14 - 6.07,  $P = 0.02$ ) groups were significant prognostic factors of PFS of mutant EGFR NSCLC.

**Conclusions:** LIPI was an independent prognostic factor of chemo-

therapy for adenocarcinoma with wild-type EGFR and of EGFR-TKI for NSCLC harboring mutant EGFR. Thus, LIPI was not a specific biomarker for ICI therapy, but a useful biomarker for chemotherapy and EGFR-TKI therapy in specific subsets of NSCLC.

**Keywords:** Lung immune prognostic index; Derived neutrophil to lymphocyte ratio; Lactate dehydrogenase; Non-small cell lung cancer; Adenocarcinoma; Squamous cell carcinoma; Epidermal growth factor receptor tyrosine kinase inhibitor; First-line cytotoxic chemotherapy

## Introduction

Non-small cell lung cancer (NSCLC) has been classified into several subsets according to histological and genetic characteristics in the past decade: squamous cell carcinoma and non-squamous NSCLC with or without driver mutations such as epidermal growth factor receptor (EGFR) mutation and anaplastic lymphoma kinase (ALK) rearrangement. Nowadays, programmed death-ligand 1 (PD-L1) expression in tumor has become a new biomarker of immune checkpoint inhibitors (ICIs) for NSCLC. Treatment strategy is various among subsets. For patients with NSCLC harboring a driver mutation, a specific tyrosine kinase inhibitor (TKI) is recommended as the first-line regimen. For squamous cell carcinoma and NSCLC without any driver mutations, either conventional cytotoxic chemotherapy or ICI is considered as the first-line regimen according to PD-L1 expression in tumor. Conventional chemotherapeutic regimen is also different between squamous and non-squamous NSCLC.

Lung immune prognostic index (LIPI) was recently developed on the basis of the combination of baseline derived neutrophil to lymphocyte ratio (dNLR) greater than 3 and lactate dehydrogenase (LDH) greater than upper limit of normal (ULN) range of each center. This index stratifies patients into three groups (good, intermediate and poor) according to the number of factors. For patients with advanced NSCLC, poor baseline LIPI was correlated with poor outcomes of overall survival (OS), progression-free survival (PFS) and disease control rate (DCR) for ICIs therapy of PD-1/PD-L1 inhibitors, but not for chemotherapy [1]. Thus, LIPI has been expressed as a specific biomarker for ICIs. The dNLR consists of easily measured hematological components of leukocyte and neutro-

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phil counts, and has similar prognostic value to the neutrophil to lymphocyte ratio (NLR) [2]. Compared with similar inflammation-based scores, dNLR has an advantage of widespread utilization, but has not been evaluated in NSCLC. Both LDH [3-5] and NLR [6-8] have been demonstrated as a useful prognostic marker for various stages and settings of patients with NSCLC. Thus, we doubted the specificity of LIPI for ICI therapy. In this study, we aimed to investigate our hypothesis that LIPI is also a prognostic marker for NSCLC patients treated with cytotoxic chemotherapy. Unlike the previous study [1], we analyzed patients according to histological and genetic subsets.

## Materials and Methods

We retrospectively collected three cohorts according to histological and genetic backgrounds: 1) Adenocarcinoma without active EGFR mutations; 2) NSCLC harboring active EGFR mutation, and 3) Squamous cell carcinoma. The patients met all the following criteria: 1) Between July 2007 and August 2017 at our hospital, first-line cytotoxic chemotherapy or EGFR-TKI monotherapy, irrespective of chemotherapeutic lines, being initiated for patients with wild-type EGFR adenocarcinoma and squamous cell carcinoma or for patients with mutant EGFR NSCLC, respectively; 2) Histologically or cytologically diagnosed with NSCLC; 3) For non-squamous NSCLC, the peptide nucleic acid-locked nucleic acid PCR clamp method or EGFR gene mutation analysis COBAS version 2 by LSI Medience Cooperation (Tokyo, Japan) confirmed wild-type or positive EGFR mutation status; 4) For adenocarcinoma with wild-type EGFR status, immunohistochemically negative or unknown anaplastic lymphoma kinase (ALK) rearrangement; 5) c-stage IIIB or IV based on the seventh tumor, node, metastasis (TNM) classification of lung cancer by the Union for International Cancer Control (UICC), or recurrence after curative-intent thoracic surgery or radiotherapy without adjuvant chemotherapy, and only for NSCLC with active EGFR mutation, c-stage IIIA was included when neither curative operation nor radiotherapy was intended; and 6) Available blood sample within 2 weeks prior to the first day of the first-line cytotoxic chemotherapy or EGFR-TKI monotherapy. We excluded patients who had received the first-line pembrolizumab, which Japanese medical insurance approved in December 2016. We collected clinical data from the electrical medical records, including age, sex, histology, EGFR mutation status, ALK rearrangement status, PD-L1 tumor proportion score, pretreatment laboratory data, chemotherapeutic regimens, its efficacy and survival. Pretreatment routine blood tests collected LDH level, total protein and albumin concentration, absolute counts of leucocyte and proportion of neutrophil in leukocyte. The dNLR was calculated by the following formula: neutrophil count/(leucocyte count - neutrophil count) [2]. The LIPI was formed by combination of dNLR > or ≤ 3 and LDH > or ≤ ULN range of each center (225 IU/L in our hospital). Patients were categorized into three groups (good, no factors; intermediate, one factor; and poor, two factors) [1].

PFS and OS were defined as the interval from the first day

of the regimen to the date of documented progressive disease (PD) or death, respectively. Based on Response Evaluation Criteria in Solid Tumors (RECIST) version 1.1 [9], we evaluated response to chemotherapy or EGFR-TKI. The data cut-off was July 31, 2018. The Osaka Police Hospital Ethics Committee approved this study. Considering the characteristic of anonymous and retrospective data, the written informed consents were waived in this study.

## Data analysis

The continuous, categorical and survival data were expressed as the median with interquartile range (IQR), frequency, median with 95% confidential intervals (CIs), respectively. Using Fisher's exact test, Kruskal-Wallis rank sum test, Kaplan-Meier method and log-rank test, we compared the relative frequencies, continuous variables, survival times of three groups, respectively. Using the Bonferroni method for the multiple comparisons, P values were corrected. In multivariate Cox proportional hazard analyses, independent variables were selected and pre-defined according to the previous study [1]. The results were described as hazard ratios (HRs) with 95% CI. P < 0.05 was defined as statistically significant. All statistical analyses were performed with EZR (Saitama Medical Center, Jichi Medical University, Saitama, Japan), which is a graphical user interface for R (The R Foundation for Statistical Computing, Vienna, Austria) [10].

## Results

We collected 175 patients with wild-type EGFR adenocarcinoma, 131 patients with NSCLC harboring active EGFR mutation and 110 patients with squamous cell carcinoma. These patients were divided into three groups by LIPI category. Tables 1-3 show patients' background, treatment, efficacy and laboratory data in each group. Table 4 shows outcomes of survival and treatment at the time of data cut-off.

In wild-type EGFR adenocarcinoma (good, intermediate and poor LIPI groups: median 19.6, 11.5 and 3.3 months, P < 0.01, respectively) and EGFR mutant NSCLC (45.4, 25.6 and 15.7 months, P < 0.01), the good LIPI group survived longer than the intermediate and poor LIPI groups (Fig. 1a, b). In contrast, in squamous cell carcinoma, there was no significant difference in OS among the three LIPI groups (13.7, 14.6 and 8.8 months, P = 0.39) (Fig. 1c). In EGFR mutant NSCLC, the PFS of good LIPI group was significantly longer than those of the other two groups (16.6, 12.6 and 8.3 months, P < 0.01) (Fig. 2b), while no significant difference was found among the three groups of wild-type EGFR adenocarcinoma (6.3, 4.3 and 2.9 months, P = 0.12) and squamous cell carcinoma (4.9, 5.9 and 3.3 months, P = 0.25) (Fig. 2a, c).

As poorer prognostic factors of OS, multivariate Cox proportional hazard analyses found intermediate group (HR 1.49, 95% CI 1.03 - 2.15, P = 0.04) of wild-type EGFR adenocarcinoma, and intermediate (HR 2.30, 95% CI 1.33 - 3.99, P < 0.01) and poor (HR 2.76, 95% CI 1.03 - 7.42, P = 0.04)

**Table 1.** Baseline Characteristics of Adenocarcinoma Without Active EGFR Mutations

	N	All	Lipi		P
			Good	Intermediate	
Backgrounds					
Sex (N), male/female	120/55	55/30	47/21	18/4	0.33 <sup>a</sup>
Age (years), median (IQR)	68 (62.4 - 74.6)	68 (63.5 - 74.9)	65 (60.2 - 74.5)	69 (64.0 - 71.5)	0.49 <sup>b</sup>
Smoking status, none/former/current smokers	30/56/89	14/26/45	13/24/31	3/6/13	0.85 <sup>a</sup>
PD-L1 status, ≥ 50%/1-49%/< < 1%/NA	85/1/161	6/4/174	2/0/0/66	0/0/0/22	0.25 <sup>a</sup>
ECOG-PS, 0 - 1/≥ 2	136/39	76/9	52/16	8/14	< 0.01 <sup>a</sup>
Stage, IIIB/IV/recurrence	32/132/11	20/57/8	12/56/0	0/19/3	< 0.01 <sup>a</sup>
Metastatic sites, ≥ 2	89	38	33	18	0.06 <sup>a</sup>
First-line chemotherapy					
Regimen					
Single/combination (N)	7/168	4/81	3/65	0/22	0.88 <sup>a</sup>
Penetraxed-containing (N)	84	40	33	11	0.96 <sup>a</sup>
Bevacizumab-containing	35	24	9	2	0.03 <sup>a</sup>
Efficacy					
ORR (%) (95% CI)	39.4 (32.1 - 47.1)	55.3 (44.1 - 66.1)	30.9 (20.2 - 43.3)	18.2 (5.2 - 40.3)	< 0.01 <sup>a</sup>
DCR (%) (95% CI)	70.9 (63.5 - 77.5)	82.4 (72.6 - 89.8)	64.7 (52.2 - 75.9)	45.5 (24.4 - 67.8)	< 0.01 <sup>a</sup>
Second and further line					
Second-line (N)	107	57	41	9	0.08 <sup>a</sup>
Immuno-checkpoint inhibitor (N)	25	18	7	0	0.02 <sup>a</sup>
Laboratory data					
dNLR					
Median (IQR)					
≥ 3 (N)	132	85	47	0	< 0.01 <sup>a</sup>
LDH					
Median (IQR)					
> ULN (N)	69	0	47	22	< 0.01 <sup>a</sup>

CI: confidence interval; DCR: disease control rate; dNLR: derived neutrophil-to-lymphocyte ratio; ECOG-PS: Eastern Cooperative Oncology Group Performance Status; IQR: interquartile range; LDH: lactate dehydrogenase; LIPI: lung immune prognostic index; NA: not assessed; ORR: overall response rate; ULN: upper limit of normal. <sup>a</sup>Fisher's exact test; <sup>b</sup>Kruskal-Wallis rank sum test.

**Table 2.** Baseline Characteristics of NSCLC With Active EGFR Mutation

	N	All	LIPI		P
			Good	Intermediate	
Backgrounds					
Sex (N), male/female	131	69	52	10	
Age (years), median (IQR)	52/79	28/41	20/32	4/6	0.96 <sup>a</sup>
Smoking status, non/former/current smokers	73 (65 - 78)	74 (67 - 78)	70.5 (63.8 - 77)	65 (50 - 81.8)	0.29 <sup>b</sup>
EGFR mutation status, Ex19el/Ex21 point/minor	63/41/27	37/21/11	22/17/13	4/3/3	0.31 <sup>a</sup>
PD-L1 status, ≥ 50%/ <sup>a</sup> 1-49%/ <sup>b</sup> < 1%/NA	129/56/7	34/31/4	27/22/3	7/3/0	0.87 <sup>a</sup>
ECOG-PS, 0 - 1≥ 2	3/4/5/119	3/1/4/61	0/3/1/48	0/0/10	0.49 <sup>a</sup>
Stage, III/IV/recurrence	93/37	55/14	37/15	2/8	< 0.01 <sup>a</sup>
Metastatic sites, ≥ 2	10/90/31	7/42/20	2/39/11	1/9/0	0.12 <sup>a</sup>
EGFR-TKI	91	43	41	7	0.15 <sup>a</sup>
Regimen, gefitinib/erlotinib/afatinib	80/38/13	43/21/5	33/13/6	4/4/2	0.42 <sup>a</sup>
Line, first/second or further	99/32	53/16	37/15	9/1	0.46 <sup>a</sup>
Efficacy					
ORR (%) (95% CI)	64.1 (55.3 - 72.3)	68.1 (55.8 - 78.8)	63.5 (49.0 - 76.4)	40.0 (12.2 - 73.8)	0.26 <sup>a</sup>
DCR (%) (95% CI)	82.4 (74.8 - 88.5)	85.5 (75.0 - 92.8)	82.7 (69.7 - 91.8)	60.0 (26.2 - 87.8)	0.16 <sup>a</sup>
Post-EGFR-TKI therapy					
Further-line	68	30	30	8	0.06 <sup>a</sup>
Osimertinib	26	9	13	4	0.06 <sup>a</sup>
Immuno-checkpoint inhibitor	5	4	1	0	0.59 <sup>a</sup>
Laboratory data					
dNLR					
Median (IQR)	1.98 (1.44 - 2.72)	1.67 (1.31 - 2.04)	2.20 (1.57 - 3.10)	3.43 (3.27 - 3.83)	< 0.01 <sup>b</sup>
≥ 3 (N)	25	0	15	10	< 0.01 <sup>a</sup>
LDH					
Median (IQR)	203 (176.5 - 240.5)	183 (163 - 198)	242 (211.8 - 308)	247 (232.5 - 466.5)	< 0.01 <sup>b</sup>
> ULN	25	0	37	10	< 0.01 <sup>a</sup>

CI: confidence interval; DCR: disease control rate; dNLR: derived neutrophil-to-lymphocyte ratio; ECOG-PS: Eastern Cooperative Oncology Group Performance Status; EGFR: epidermal growth factor receptor; IQR: interquartile range; LDH: lactate dehydrogenase; LIPI: lung immune prognostic index; NA: not assessed; ORR: overall response rate; TKI: tyrosine kinase inhibitor; ULN: upper limit of normal. <sup>a</sup>Fisher's exact test; <sup>b</sup>Kruskal-Wallis rank sum test.

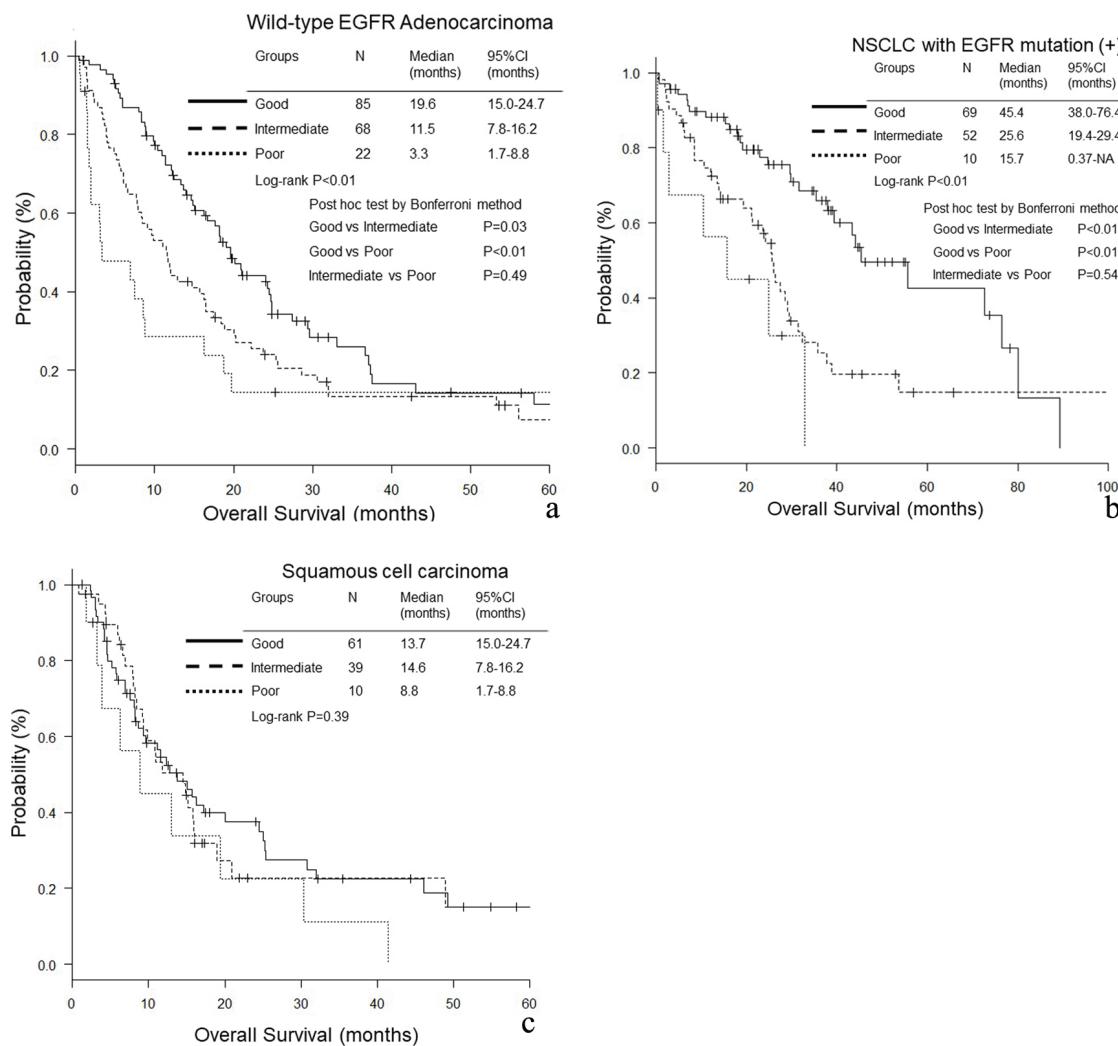
**Table 3.** Baseline Characteristics of Squamous Cell Carcinoma

	All	LIPi		P
		Good	Intermediate	
N				
Backgrounds	110	61	39	10
Sex (N), male/female	85/25	49/12	28/11	8/2
Age (years), median (IQR)	71.5 (65 - 76)	72 (65 - 76)	70 (66 - 75)	71.5 (62 - 75)
Smoking status, non/former /current smoker	8/42/60	4/19/38	3/20/16	1/3/6
PD-L1 status, ≥ 50% /1-49% /< 1%/NA	1/1/2/106	1/0/1/59	0/1/1/37	0/0/0/22
ECOG-PS, 0-1/≥ 2	83/27	50/11	28/11	5/5
Stage, IIB/IV/recurrence	40/57/13	25/25/11	14/23/2	1/9/0
Metastatic sites, ≥ 2	33	16	13	4
First-line chemotherapy				
Regimen				
Single/combination (N)	11/89	8/53	2/37	1/9
PTX or nab-PTX	72	36	28	8
Efficacy				
ORR (%) (95% CI)	42.7 (33.3 - 52.5)	45.9 (33.1 - 59.2)	35.9 (21.2 - 52.8)	50.0 (18.7 - 81.3)
DCR (%) (95% CI)	65.5 (55.8 - 74.3)	63.9 (50.6 - 75.8)	69.2 (52.4 - 83.0)	60.0 (26.2 - 87.8)
Second and further line				
Second-line (N)	66	39	23	4
Immuno-checkpoint inhibitor (N)	10	5	5	0
Laboratory data				
dNLR				
Median (IQR)	2.25 (1.73 - 3.13)	2.00 (1.69 - 2.39)	3.20 (1.88 - 4.04)	3.39 (3.19 - 3.91)
≥ 3 (N)	30	0	20	10
LDH				
Median (IQR)	198.5 (168 - 229)	181 (158 - 202)	224 (189.5 - 277.5)	282 (243 - 304.5)
> ULN	29	0	19	10

CI: confidence interval; DCR: disease control rate; dNLR: derived neutrophil-to-lymphocyte ratio; ECOG-PS: Eastern Cooperative Oncology Group Performance Status; IQR: interquartile range; LDH: lactate dehydrogenase; LIPi: lung immune prognostic index; NA: not assessed; nab-PTX: nanoparticle albumin-bound paclitaxel; ORR: overall response rate; PTX: paclitaxel; ULN: upper limit of normal. <sup>a</sup>Fisher's exact test; <sup>b</sup>Kruskal-Wallis rank sum test.

**Table 4.** Outcomes

	N	Ad, EGFR mt (-)	NSCLC, EGFR mt (+)	SQ
Survival		175	131	110
Dead		137	71	79
At our hospital		108	38	61
At other institutions		15	23	11
At home		14	10	7
Alive		20	37	10
Lost follow-up		18	23	21
Chemotherapy				
Confirmed PD or death		154	103	88
Continued		2	17	0
Discontinued		173	86	110
PD		89	73	36
Completion of pre-defined courses		21	0	27
Adverse effects		29	13	18
Deteriorated conditions		25	12	24
Refusal		8	4	3
Transfer to other hospitals		1	5	0
Unknown cause		0	3	2



**Figure 1.** Overall survival (OS) according to lung immune prognostic index (LIPI) groups. (a) Adenocarcinoma without any driver mutations. (b) NSCLC with EGFR mutation. (c) Squamous cell carcinoma.

LIPI groups of EGFR mutant NSCLC (Table 5). Multivariate analyses detected intermediate (HR 1.57, 95% CI 1.01 - 2.44, P = 0.04) and poor (HR 2.63, 95% CI 1.14 - 6.07, P = 0.02) LIPI groups as poor prognostic factors of PFS of EGFR mutant NSCLC (Table 6).

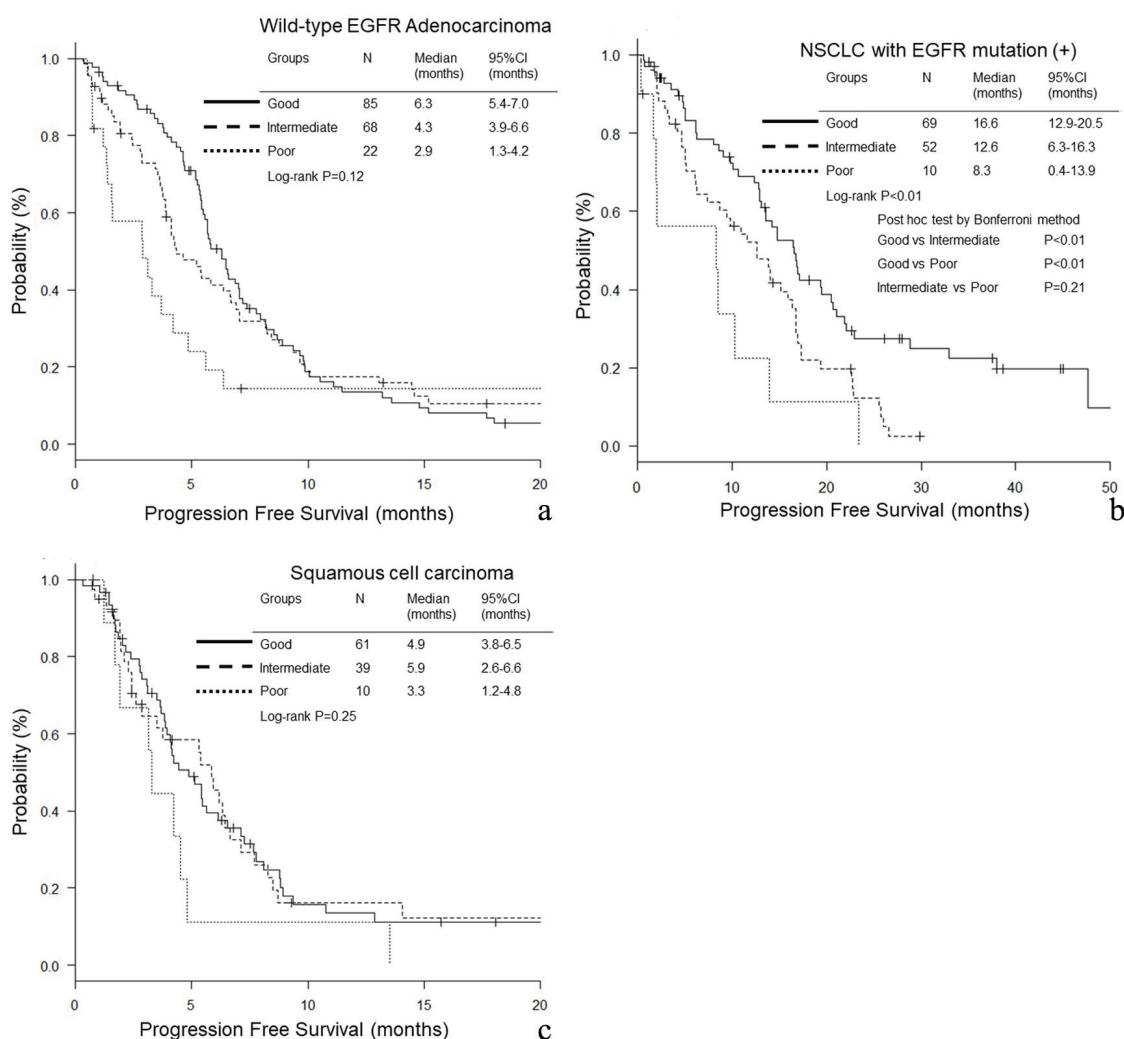
## Discussion

This study demonstrated that pretreatment LIPI is a prognostic marker for adenocarcinoma with wild-type EGFR and for NSCLC with active EGFR mutation, but not for squamous cell carcinoma. LIPI showed different trends as a prognostic marker according to histological and genetic subsets.

Our multivariate analyses and comparisons of OS and PFS showed this marker as independent prognostic factors of OS of first-line cytotoxic chemotherapy for adenocarcinoma with wild-type EGFR, and of OS and PFS of EGFR-TKI therapy for

NSCLC with active EGFR mutation. Our results were contrary to the previous study, in which LIPI was significantly correlated with outcomes for ICI, but not for chemotherapy [1]. In the study by Mezquita, chemotherapy cohort (N = 162) was composed only of patients who had received second or later chemotherapy, and included various subsets of NSCLC: 72% of adenocarcinoma, 14% of squamous cell carcinoma, 23% of positive EGFR mutation or ALK rearrangement [1]. Our study investigated first-line chemotherapy or first EGFR-TKI therapy, and divided patients by subsets. Thus, LIPI is not only a specific marker for ICI therapy, but also is a useful prognostic marker for some specific subsets of NSCLC.

In our study, LIPI was not an independent prognostic factor of chemotherapy for squamous cell carcinoma. There was a different trend of LIPI as a prognostic factor between squamous cell carcinoma and other subsets. The difference of hematological markers according to subsets was observed in our series of previous studies. As a prognostic factor of OS,



**Figure 2.** Progression-free survival (PFS) according to lung immune prognostic index (LIPI) groups. (a) Adenocarcinoma without any driver mutations. (b) NSCLC with EGFR mutation. (c) Squamous cell carcinoma.

neither NLR nor modified Glasgow prognostic score (mGPS) was useful for squamous cell carcinoma [11], while mGPS, but not NLR, was selected for adenocarcinoma with wild-type EGFR [12]. On the other hand, NLR was significantly useful for NSCLC harboring mutant EGFR, though mGPS was not investigated in our previous study [13]. Little has been reported on mGPS as a prognostic factor for patients with driver mutations. Thus, hematological biomarkers variously responded among subsets of NSCLC. It may be necessary to evaluate biomarkers by histological and genetic subsets.

Our study had some limitations. First, we are afraid of case bias owing to our retrospective, single-centered and small sample-sized study. Second, in terms of recent fast advance in NSCLC treatment, the regimens that our patients received are becoming outdated. Our study included many patients who had initiated their treatment and become unable to proceed to the further lines until the approval of ICIs or third-generation EGFR-TKI. The new era of next generation treatment is com-

ing, and it needs more cases to confirm LIPI as a useful marker of advanced NSCLC.

## Conclusions

LIPI was an independent prognostic factor of chemotherapy for adenocarcinoma with wild-type EGFR and of EGFR-TKI for NSCLC harboring mutant EGFR. Thus, LIPI was not a specific biomarker for ICI therapy, but a useful biomarker for chemotherapy and EGFR-TKI therapy in specific subsets of NSCLC. The optimal prognostic markers may be different among subsets of NSCLC.

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**Table 5.** Multivariate Analysis of Overall Survival

Variable	Ad, EGFR mt (-), HR (95% CI)	P	NSCLC, EGFR mt (+), HR (95% CI)	P	SQ, HR (95% CI)	P
Age, years						
< 70	1 (Reference)		1 (Reference)		1 (Reference)	
≥ 70	1.29 (0.90 - 1.85)	0.16	1.95 (1.13 - 3.35)	0.02	0.90 (0.56 - 1.45)	0.66
Smoking history						
Non-smoker	1 (Reference)		1 (Reference)		1 (Reference)	
Smoker	1.41 (0.88 - 2.28)	0.16	1.07 (0.64 - 1.79)	0.81	1.87 (0.57 - 6.20)	0.30
No. of metastatic sites						
< 2	1 (Reference)		1 (Reference)		1 (Reference)	
≥ 2	0.88 (0.62 - 1.26)	0.49	2.30 (1.26 - 4.22)	< 0.01	1.51 (0.90 - 2.54)	0.12
ECOG-PS						
0 or 1	1 (Reference)		1 (Reference)		1 (Reference)	
≥ 2	3.12 (2.04 - 4.79)	< 0.01	2.54 (1.33 - 4.86)	< 0.01	3.14 (1.87 - 5.30)	< 0.01
Line of EGFR-TKI						
First-line			1 (Reference)		1 (Reference)	
Second or later line			0.71 (0.40 - 1.25)		0.24	
Albumin level (g/dL)						
≥ 3.5	1 (Reference)		1 (Reference)		1 (Reference)	
< 3.5	1.56 (1.05 - 2.32)	0.03	0.59 (0.31 - 1.10)	0.10	1.17 (0.71 - 1.93)	0.54
LPI						
Good	1 (Reference)		1 (Reference)		1 (Reference)	
Intermediate	1.49 (1.03 - 2.15)	0.04	2.30 (1.33 - 3.99)	< 0.01	1.01 (0.61 - 1.67)	0.98
Poor	1.67 (0.94 - 2.98)	0.08	2.76 (1.03 - 7.42)	0.04	1.64 (0.78 - 3.44)	0.19

Ad: adenocarcinoma, CI: confidence interval, ECOG-PS: Eastern Cooperative Oncology Group Performance Status; EGFR: epidermal growth factor receptor; HR: hazard ratio; LPI: lung immune prognostic index; nab-PTX: nanoparticle albumin-bound paclitaxel; NSCLC: non-small cell carcinoma; SQ: squamous cell carcinoma; TKI: tyrosine kinase inhibitor.

**Table 6.** Multivariate Analysis of Progression-Free Survival

Variable	Ad, EGFR mt (-)		NSCLC, EGFR mt (+)		P	SQ HR (95% CI)	P
	HR (95% CI)	P	HR (95% CI)	P			
Age, years							
< 70	1 (Reference)		1 (Reference)				
≥ 70	1.09 (0.77 - 1.53)	0.62	1.25 (0.82 - 1.90)	0.30	1.13 (0.71 - 1.80)	0.61	
Smoking history							
Non-smoker	1 (Reference)		1 (Reference)				
Smoker	1.01 (0.65 - 1.58)	0.96	1.08 (0.72 - 1.60)	0.72	1.22 (0.53 - 2.82)	0.64	
No. of metastatic sites							
< 2	1 (Reference)		1 (Reference)				
≥ 2	0.99 (0.71 - 1.38)	0.94	1.52 (0.94 - 2.45)	0.09	1.10 (0.66 - 1.85)	0.72	
ECOG-PS							
0 or 1	1 (Reference)		1 (Reference)				
≥ 2	1.90 (1.26 - 2.86)	< 0.01	1.48 (0.86 - 2.55)	0.16	1.82 (1.04 - 3.18)	0.04	
Bevacizumab-containing							
Yes	1 (Reference)		1 (Reference)				
No	1.40 (0.93 - 2.12)	0.11					
PTX or nab-PTX-containing							
Yes	1 (Reference)		1 (Reference)				
No					1.53 (0.95 - 2.45)	0.08	
EGFR-TKI line							
First	1 (Reference)		1 (Reference)				
Second or later			0.76 (0.47 - 1.21)		0.24		
Albumin level, g/dL							
≥ 3.5	1 (Reference)		1 (Reference)				
< 3.5	1.15 (0.79 - 1.67)	0.48	0.80 (0.46 - 1.37)	0.41	0.90 (0.57 - 1.44)	0.67	
LIPI							
Good	1 (Reference)		1 (Reference)				
Intermediate	1.00 (0.70 - 1.43)	1.00	1.57 (1.01 - 2.44)	0.04	1.08 (0.67 - 1.73)	0.75	
Poor	1.35 (0.78 - 2.34)	0.29	2.63 (1.14 - 6.07)	0.02	2.00 (0.94 - 4.22)	0.07	

Ad: adenocarcinoma; CI: confidence interval; ECOG-PS: Eastern Cooperative Oncology Group Performance Status; EGFR-TKI: epidermal growth factor receptor-tyrosine kinase inhibitor; HR: hazard ratio; LIPI: lung immune prognostic index; nab-PTX: nanoparticle albumin-bound paclitaxel; NSCLC: non-small cell carcinoma; PTX: paclitaxel; SQ: squamous cell carcinoma.

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## Conflict of Interest

The authors declare that they have no conflict of interest.

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