Estimating Survival in Patients with Non-Small-Cell Lung Cancer and Brain Metastases: A Verification of the Graded Prognostic Assessment for Lung Cancer Using Molecular Markers (Lung-molGPA)

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Purpose: A new tool based on clinical characteristics and molecular factors (Lung-molGPA) was developed to predict the survival of patients with non-small-cell lung cancer but has not been validated. This study aims to validate the feasibility of the Lung-molGPA in NSCLC.

Patients and Methods: Patients diagnosed NSCLC between Feb 2012 and July 2018 were retrospectively reviewed and scored using the Lung-molGPA tool to compare clinical outcomes. Hazard ratios (HRs) and 95% confidence intervals (CIs) were calculated by Cox regression analyses.

Results: A total of 618 patients (524 adenocarcinoma [ADC], 94 non-adenocarcinoma [non-ADC]) were collected. For all patients, the median survival time (MST) was 33.0 months (33.6 and 28 months in the ADC and non-ADC groups, respectively; $p = 0.21$). In the ADC group, the MST for patients with a Lung-molGPA score of 3.5 to 4 was more than 4 years, while the MST was only 25 months in patients scoring 0–1, 30.0 months in patients scoring 1.5–2, and 35.0 months for scores of 2.5–3 ($p = 0.048$). For the non-ADC group, the MST for scores 0–1, 1.5–2, 2.5–3, and 3.5–4 were 12.0, 20.2, 29.0, and 33.0 months, respectively ($p = 0.017$).

Conclusion: Our findings provided evidence validating the Lung-molGPA score as a useful tool to determine treatment strategies and to predict prognosis. The model is still exploratory and needs to be evaluated further in combination with additional prognostic markers.

Keywords: NSCLC, GPA, brain metastasis, survival, prognosis

Introduction

Lung cancer is the first leading cause of cancer-related deaths in China and worldwide.^{1,2} Brain metastases (BM) are frequently observed in non-small-cell lung cancer (NSCLC), which undoubtedly exerts a devastating influence on the survival and quality of life of patients. About 40% of patients present BM during the course of the disease.^{3,4} The median overall survival (OS) was reported to be merely 4–6 months after whole brain radiation therapy (WBRT) in the chemotherapy era.^{3,4}

With the recent advent of molecularly targeted therapies, it is evident that NSCLC patients with BM are an extremely heterogeneous population. Identifying various subgroups with different prognosis is essential to precisely individualize treatment and especially to design clinical trials. Several models have

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focused on predicting survival for patients with BM such as Recursive Partitioning Analysis (RPA), which was established by the Radiation Therapy Oncology Group (RTOG) 20 years ago, based on the data from 1200 patients in 3 clinical trials.⁵ In this model, patients are stratified according to age, the Karnofsky Performance Status (KPS), control of primary tumor, and extracranial metastases, into three classes of disease with median survival ranging from 2.3 to 7.1 months. Furthermore, the Diagnosis Specific Graded Prognostic Assessment (DS-GPA) score was established in 2012 and takes into consideration age, and four clinical factors: KPS, extracranial metastases, and the number of brain metastases.⁶ Median survival ranges from 3.0 to 14.8 months according to the four classes of disease.

It is known that advanced NSCLC patients harboring epidermal growth factor receptor (*EGFR*) and anaplastic lymphoma kinase (*ALK*) alterations treated with tyrosine kinase inhibitors (TKIs), achieve an OS superior to that of patients without any driving gene mutations. However, NSCLC patients with *EGFR* mutations experience a higher incidence of BM, ranging from 44% to 63%. Thus, to predict the survival of NSCLC with BM, molecular factors should also be considered. Sperduto et al updated the graded prognostic assessment for Lung Cancer Using Molecular Markers (Lung-molGPA) based on the clinical characteristics of 2186 patients to include age, KPS, extracranial metastases, number of BM, and *EGFR* and *ALK* alterations in patients with ADC.⁷ The median OS for the entire cohort was 12 months, and patients harboring *EGFR* and *ALK* alterations and with Lung-molGPA scores of 3.5 to 4.0 had a median survival of nearly 4 years.

Nonetheless, published studies providing evidence for the validation of the Lung-molGPA based on large patient samples are lacking. Hence, we conducted the present study and reviewed the records of 618 patients, with the aim to validate the clinical utility of the Lung-molGPA scoring system.

Patients and Methods

Patients

The records of patients diagnosed with NSCLC treated for BM between February 2012 and June 2018 at our cancer center were reviewed. All patients were staged according to the criteria of AJCC 7th edition. At the initial diagnosis, whole-body systemic evaluation was performed to evaluate the disease stage, including history and physical examination, blood profile, cervical ultrasound or computed tomography

(CT), chest and abdomen enhanced contrast CT, brain enhanced contrast CT or magnetic resonance imaging, positron emission tomography-CT was not routinely performed in our cancer center and data was available only for a few patients.

Graded Prognostic Assessment

All patients were stratified based on the Lung-molGPA score. The Lung-molGPA score was calculated according to the graded prognostic assessment (GPA) index using a web-based interface (<http://brainmetgpa.com/>), and the supplementary calculations were carried out according to those reported in previous studies.⁷ The type of treatment was not considered because the purpose of the Lung-molGPA score is to estimate survival prior to treatment. In the Lung-molGPA scoring system, according to the size of the effect on the survival, KPS from 90 to 100, no extracranial metastases, >4 BM and *EGFR* or *ALK* positivity were given a score of 1, age <70 years and 1–4 BM were given a score of 0.5. Thus, the maximum score remained 4.0. The parameters of the new Lung-molGPA are detailed in Table 1. All enrolled patients were divided into four groups based on the final scores: 0–1, 1.5–2, 2.5–3, 4 (Table 1).

Statistical Analysis

OS was defined from the date of initial treatment of BM to the date of death due to any cause, or to the latest follow-up. Progression-free survival (PFS) was defined as the time from the date of initial treatment of BM to the date of disease progression, or death due to any cause. OS and PFS were calculated by the Kaplan-Meier method, and differences in survival curves between the groups were evaluated by the Log rank test. Two-sided *p* values < 0.05 were considered

Table 1 Lung-molGPA Scoring Chart for NSCLC

Parameter	GPA Scoring Criteria		
	0	0.5	1
Age, yrs	≥ 70	< 70	NA
KPS	< 80	80	90–100
ECM	Present		Absent
BM	> 4	1–4	NA
Gene Status	<i>EGFR</i> neg/unk and <i>ALK</i> neg/unk	NA	<i>EGFR</i> pos or <i>ALK</i> pos

Abbreviations: GPA, graded prognostic assessment; KPS, Karnofsky Performance Status; ECM, extracranial metastases; BM, brain metastases; NA, not applicable; neg/unk, negative or unknown; NSCLC, non-small-cell lung cancer; pos, positive; *EGFR*, epidermal growth factor receptor; *ALK*, anaplastic lymphoma kinase.

statistically significant. Hazard ratios (HRs) and 95% confidence intervals (CIs) were calculated by Cox regression analyses.

Results

Patient Characteristics

During the period from February 2012 to June 2018, the records of 618 patients diagnosed with NSCLC and BM were enrolled in this retrospective study. A total of 524 patients were diagnosed with adenocarcinoma (ADC) and the remaining 94 patients were diagnosed with non-adenocarcinoma (non-ADC), which includes squamous cell lung cancer (42 patients), large cell lung cancer (20 patients) and cancers (32 patients). Because driver gene detection is not covered by government health insurance, only 352 (57.0%) of patients' driver genes were determined, with 333 patients in the ADC group and 19 patients in the non-ADC. *EGFR* or *ALK*-positive lesions were found in 224 (67.3%) patients, with 218 patients in the ADC group and 6 patients in the non-ADC group. At the initial diagnosis, 240 patients presented with symptoms of BM, including 200 patients in

the ADC group and 40 patients in the non-ADC group. The patient characteristics are detailed in [Tables 2](#) and [3](#).

According to the Lung-molGPA score criteria detailed in [Table 1](#), each patient in the group was given a score. As a result, 38.5% of patients (238/618) and 45.3% of patients (280/618) scored 1.5–2 and 2.5–3, respectively, whereas only 7% (43/618) and 10.8% (67/618) scored 0–1 and 3.5–4, respectively. In the ADC subgroup, 37 (7.1%) patients scored 0–1, 201 (38.4%) patients scored 1.5–2, 234 (44.6%) patients scored 2.5–3, and 52 (9.9%) patients in score 3.5–4 group, whereas was 6 (6.4%) patients in score 0–1 group, 37 (39.4%) patients in score 1.5–2 group, 36 (38.3%) patients in score 2.5–3 group and 15 (15.9%) patients in score 3.5–4 group, in the non-ADC group, respectively ([Table 4](#)).

Treatment Summary

First-Line Systemic Treatment

Because the TKIs were not covered by the government health insurance before the year 2017 in China, only 118 patients (52.6%) were given TKIs among the 224 patients harboring *EGFR*+ or *ALK*+, and all were first-generation including

Table 2 Univariate and Multivariate Cox Regression Analyses for OS and PFS in Adenocarcinoma Patients

Parameter	Categories	N	Univariate		Multivariate	
			OS	PFS	OS	PFS
			HR (95% CI)	HR (95% CI)	HR (95% CI)	HR (95% CI)
KPS	90–100	210	I	I	I	I
	80	263	1.59 (1.07–2.36)	1.17 (0.91–1.49)	1.35 (0.89–2.04)	1.10 (0.71–1.69)
	< 80	51	1.34 (0.69–2.62)	1.04 (0.68–1.58)	1.19 (0.60–2.35)	1.16 (0.76–1.77)
Age	<70	459	I	I	I	I
	≥70	65	2.40 (1.49–3.85)	1.92 (1.36–2.69)	2.20 (1.34–3.60)	1.97 (1.38–2.81)
ECM	Absent	154	I	I	I	I
	Present	370	0.93 (0.63–1.37)	1.32 (1.02–1.71)	1.06 (0.71–1.59)	1.45 (1.10–1.90)
BM	I–4	325	I	I	I	I
	>4	184	1.51 (0.47–4.84)	2.28 (0.73–7.16)	1.43 (0.44–4.72)	2.36 (0.75–7.42)
	NA	15	0.98 (0.67–1.42)	1.82 (0.58–5.74)	0.97 (0.66–1.42)	2.03 (0.64–6.45)
Gene Status	<i>EGFR</i> pos or <i>ALK</i> pos	218	I	I	I	I
	<i>EGFR</i> neg/unk and <i>ALK</i> neg/unk	115	2.31 (1.44–3.70)	1.47 (1.10–1.98)	2.17 (1.33–3.54)	1.49 (1.10–2.02)
	NA	191	1.56 (1.01–2.41)	0.99 (0.75–1.30)	1.59 (1.33–3.54)	1.09 (0.83–1.44)
Symptom	No	324	I	I	I	I
	Yes	200	1.08 (0.73–1.58)	1.32 (1.02–1.71)	1.06 (0.71–1.59)	1.45 (1.10–1.90)

Abbreviations: OS, overall survival; PFS, progression-free survival; KPS, Karnofsky Performance Status; ECM, extracranial metastases; BM, brain metastases.

Table 3 Univariate and Multivariate Cox Regression Analyses for OS and PFS in Nonadenocarcinoma Patients

Parameter	Categories	N	Univariate		Multivariate	
			OS	PFS	OS	PFS
			HR (95% CI)	HR (95% CI)	HR (95% CI)	HR (95% CI)
KPS	90–100	35	I	I	I	I
	80	47	1.79 (0.72–4.45)	0.71 (0.29–1.73)	2.10 (0.79–5.55)	0.53 (0.18–1.52)
	≤70	12	2.57 (0.88–7.49)	1.14 (0.49–2.67)	2.93 (0.71–12.02)	1.04 (0.39–2.74)
Age	<70	79	I	I	I	I
	≥70	15	0.76 (0.23–2.54)	1.30 (0.55–3.06)	0.68 (0.19–2.49)	1.51 (0.60–3.79)
ECM	Absent	45	I	I	I	I
	Present	49	1.45 (0.65–3.23)	1.03 (0.59–1.81)	2.40 (0.90–6.39)	1.17 (0.62–2.20)
BM	1–4	66	I	I	I	I
	>4	28	2.61 (1.20–5.67)	1.08 (0.60–1.98)	2.60 (0.96–7.01)	1.12 (0.54–2.32)
Gene Status	EGFR pos or ALK pos	6	I	I	I	I
	EGFR neg/unk and ALK neg/unk	13	2.01 (0.47–8.57)	2.20 (0.69–7.05)	1.94 (0.44–8.52)	3.67 (0.99–13.70)
	NA	75	0.60 (0.05–6.59)	1.21 (0.43–3.44)	0.11 (0.01–1.75)	1.75 (0.56–5.50)
Symptom	No	54	I	I	I	I
	Yes	40	1.12 (0.52–2.46)	1.07 (0.60–1.89)	1.84 (0.73–4.63)	0.89 (0.46–1.71)

Abbreviations: OS, overall survival; PFS, progression-free survival; KPS, Karnofsky Performance Status; ECM, extracranial metastases; BM, brain metastases.

Table 4 Comparison Survival in Patients with NSCLC

Lung GPA Score	Adenocarcinoma NSCLC Lung-molGPA		Nonadenocarcinoma NSCLC Lung-molGPA	
	MS, Mo	Patients, No. (%)	MS, Mo	Patients, No. (%)
0–1	25	37 (7.06%)	12	6 (6.38%)
1.5–2	30	201 (38.36%)	20.2	37 (39.36%)
2.5–3	35	234 (44.66%)	29	36 (38.30%)
3.5–4	NA	52 (9.92%)	33	15 (15.96%)
Overall	33.6	524 (100%)	28	94 (100%)

Abbreviations: GPA, graded prognostic assessment; MS, median survival; NSCLC, non-small-cell lung cancer.

gefitinib, erlotinib, and icotinib. At the last follow-up, 97 patients had experienced disease progression, and the second biopsy was conducted in 75 patients (77.3%). As a result, 40 patients (53.3%) harbored T790M mutation, and of these, 32 patients received the third-generation TKI, osimertinib.

The remaining 500 patients were given platinum-based chemotherapy as the first-line treatment, which included

pemetrexed, paclitaxel, and gemcitabine. In the 106 patients harboring *EGFR*+, 80 patients were given TKI as the second-line treatment when the patients experienced suffered from disease progression.

Local Therapy to Brain Metastasis

The local treatment for the 391 patients with 1–4 BM was as follows: WBRT ± boost (or simultaneous integrated boost) to 135 patients (34.5%); SRS (or SBRT) alone to 202 patients (51.7%); and surgery alone to 54 patients (13.8%).

For 212 patients with more than 4 BM, 182 patients (85.8%) received WBRT, while 30 patients (14.2%) received SBRT alone. For the 15 patients with not applicable information, WBRT was conducted in 12 patients and WBRT+SRS in 3 patients.

Overall Survival

The last follow-up was 31 August, 2019. The median follow-up time of the whole group was 26.8 months. The median survival time (MST) of the whole group and the OS rate at 3 years were 33.0 months and 44.5%, respectively. For patients with ADC, the MST and the OS rates at

Figure 1 Survival curves of patients with ADC and non-ADC.

3 years were 33.6 months and 45.6%, which were higher than the 28 months and 37.2% observed in the subgroup with non-ADC but the difference was not significant ($p = 0.21$). The survival curves are compared in [Figure 1](#) and in [Table 4](#).

Impacts of Clinical Characteristics on OS and PFS

For ADC patients, on univariate analysis, older age, lower KPS, presence of ECM, additional brain metastatic lesions, *EGFR* negative/unknown, and *ALK* negative/unknown were significantly associated with increased risk of death and disease progression of ADC patients with BM ($HR > 1$). As we can see in [Table 2](#), multivariate analysis demonstrated that older age, worse performance status, presence of ECM, and local brain treatment were significantly associated with an increased risk of death and disease progression of ADC patients with BM ($HR > 1$). Furthermore, univariate and multivariate Cox regression analyses showed the same results in terms of prognostic factors for non-ADC patients.

Survival Stratified by Lung-molGPA for NSCLC

For ADC patients, the survival rates of patients stratified according to the four prognostic classes are detailed in [Table 3](#). For 52 patients with Lung-molGPA scores of 3.5 to 4.0, the MST was not reached. However, this group had a median survival of more than 4 years. The survival difference between the four score classes was significant ($p = 0.048$) ([Figure 2](#)). For non-ADC patients,

Figure 2 Survival curves stratifying lung adenocarcinoma patients by the Lung-molGPA.

only 15 patients had Lung-molGPA scores of 3.5 to 4.0 ($n = 15$), the MST in this subgroup was 33 months. The survival difference between the four score classes was significant ($p = 0.017$) ([Figure 3](#)).

For ADC patients, there was no statistically significant difference across groups stratified by the Lung-molGPA scores ($p = 0.437$ for 0–1 vs 1.5–2; $p = 0.072$ for 1.5–2 vs 2.5–3; and $p = 0.304$ for 2.5–3 vs 3.5–4). However, the MST in the score 1.5–2 group was significantly lower than that in the score 3.5–4 group (30 months vs not reached, $p = 0.049$).

Similarly, no statistically significant difference in survival was observed for groups stratified by the Lung-molGPA in non-ADC patients ($p = 0.08$ for 0–1 vs 1.5–2; $p = 0.26$ for 1.5–2 vs 2.5–3; $p = 0.38$ for 2.5–3 vs

Figure 3 Survival curves stratifying lung non-adenocarcinoma patients by the Lung-molGPA.

3.5–4). However, the survival in patients scoring 0–1 was significantly lower than that in patients scoring 2.5–3 or 3.5–4 ($p = 0.02$ and $p = 0.005$, respectively).

PFS in ADC patients scoring 3.5–4 was significantly longer than in patients with scores of 0–1 or 1.5–2 (17 months vs 9 months, $p = 0.01$; 17 months vs 10 months, $p = 0.022$; respectively; [Figure S1](#)). In addition, the PFS with score 0–1 was significantly lower than that with score 1.5–2 ($p = 0.041$; [Figure S2](#)). As we can see in [Table S1](#), we found that the baseline clinical features were matched between the two groups, however, more EGFR/ALK-positive patients in adenocarcinoma group compared with nonadenocarcinoma group. These suggest that adenocarcinoma patients have more treatment options such as TKIs, so adenocarcinoma patients have a longer survival and better prognosis.

Discussion

BM are a serious risk factor for poor survival in patients with NSCLC. Emerging evidence has revealed that patients with BM could benefit from targeted therapy and immunotherapy.^{8,9} Therefore, we identified patient subgroups stratified by survival outcomes based on the patient's intrinsic characteristics that were associated with the combination of treatment strategies. Patients wild type for mutations aside from radiotherapy underwent treatment with mono-chemotherapy according to guidelines for elderly people.²⁷ RPA, GPA, and DS-GPA have been validated to predict survival,^{10,11} but they no longer meet the clinical needs given the availability of new treatment strategies. Sperduto et al proposed a Lung-molGPA model, which was based on age, KPS, extracranial metastases, number of BM, EGFR and ALK alterations, to stratify patients with different survival outcomes.⁷ The final results showed the median survival for ADC patients with Lung-molGPA scores of 0–1, 1.5–2, 2.5–3, and 3.5–4 were 6.9 months, 13.7 months, 26.5 months, and 46.8 months, respectively ($p = 0.03$ for 0–1 vs 1.5–2; $p < 0.001$ for 1.5–2 vs 2.5–3; and $p < 0.001$ for 2.5–3 vs 3.5–4.0). Similarly, the survival of non-ADC patients with lung GPA scores 0–1, 1.5–2, and 2.5–3 were 5.3 months, 9.8 months, and 12.8 months, respectively ($p < 0.001$ for 0–1 vs 1.5–2; and $p = 0.04$ for 1.5–2 vs 2.5–3). However, no studies focusing on the Lung-molGPA model have been performed to validate their clinical utility since then.

In the present study, we retrospectively reviewed 618 Asian patients newly diagnosed NSCLC with BM to evaluate the utility of the Lung-molGPA model. The MST in

the ADC group was prolonged as the Lung-molGPA score increased; however, there was no significant difference in each of the groups. Notably, the MST for patients with score 3.5–4 was more than 4 years, which was significantly longer than that for patients with score 1.5–2 (30 months, $p = 0.049$). Moreover, PFS in ADC patients with a score of 3.5–4 was significantly longer than patients having scores 0–1 or 1.5–2 (17 months vs 9 months, $p = 0.01$; 17 months vs 10 months, $p = 0.022$; respectively; [Figure S1 new](#)). This study was the first to consider the effects of driver genes in non-ADC and to investigate its impact on stratification of patient outcomes. Nineteen patients (19/94, 20.2%) underwent genetic testing of driver genes, of whom 6 patients (31.5%) were identified with EGFR or ALK mutations. OS with scores 0–1, 1.5–2, 2.5–3, and 3.5–4 according to the Lung-molGPA grade was 12 months, 20.2 months, 29 months, and 33 months, in the non-ADC group, respectively, and OS was not significantly different across groups. Nonetheless, the worst survival was observed in patients who scored 0–1, which was significantly lower than patients who scored 2.5–3 and 3.5–4 ($p = 0.02$ and $p = 0.005$, respectively). In addition, the PFS observed in patients with a score of 0–1 was significantly lower than in those with scores of 1.5–2 ($p = 0.041$; [Figure S2 new](#)).

The Lung-molGPA, as a new prognostic tool, provides an accurate estimation of prognosis for NSCLC patients with BM.⁷ Patient age, KPS, presence of extracranial metastases, number of BM, and positive findings for EGFR and ALK were added to the Lung-molGPA.^{12,13} In this study, consistent with previously published data,^{14–16} age, Karnofsky Performance Status, extracranial metastases, number of BM, and gene status, were all risk factors for OS in both ADC and non-ADC patients.²⁸ The Lung-molGPA was verified in NSCLC patients with BM. Although there was no statistically significant difference between the four groups, there were significant differences in median survival and survival rates. In the present study, NSCLC patients with BM showed a median OS of 11 months, which was slightly longer than the median OS of 8.6 months reported in Hendriks et al's study.¹⁷ This survival advantage might be associated with the widespread use of molecularly-targeted agents as a treatment strategy. Moreover, additional treatment options after disease progression may have also contributed to improved survival.

An excellent clinical tool should have the ability to guide the choice of clinical treatment strategies. A secondary analysis of RTOG 95-08 indicated

a survival advantage regardless of the patient had 1, 2, or 3 BM for patients with a high GPA (3.5–4).¹⁸ This result strengthens the observations that SRS, when given in association with WBRT, improves local disease control and OS in those patients with optimal prognostic factors. Another retrospective study, including 164 patients treated with SRS alone or SRS plus WBRT, indicated that the MST stratified by DS-GPA was consistent with the OS associated with BM in NSCLC DS-GPA and did not correlate with SRS plus WBRT or SRS alone.¹⁹ According to the current evidence, only patients with specific characteristics (such as no more than 4 BM) should be considered for SRS.^{20,21} Nagtegaal et al conducted a retrospective study to assess the feasibility of DS-GPA in 367 NSCLC patients treated with SRS for BM.²² Their findings indicated that the median survival was significantly shorter in the ADC NSCLC subgroup with GPA scores of 2.5–3 (15.4 months vs 26.5 months), but in other groups, it was largely comparable to the corresponding DS-GPA cohorts. The difference may be due to the specific characteristics of patients treated with SRS versus other patients. Hence, the value of DS-GPA as a useful prognostic tool was confirmed for counseling of individual patients with BM before undergoing SRS. Furthermore, for patients treated with WBRT, the MST for GPA 0–1 with elevated serum lactate dehydrogenase (LDH) was only 37 days compared to 123 days in patients without elevated LDH ($p < 0.001$).²³ Hence, the stratification of prognosis according to GPA before treatment is necessary and contributes to the choice of treatment. Conversely, targeted therapy as the first-line treatment for patients with *EGFR* mutations with BM is now widely recognized. In addition, earlier guidelines recommended treatment with *EGFR* TKIs was suitable for locally advanced or metastatic NSCLC after the failure of the first-line chemotherapy.^{24,25}

In summary, if the patient presents with 1 to 3 BM with a high-grade GPA (3.5–4), SRS can be given combined with WBRT; for patients with *EGFR* mutations and BM, targeted therapy is an optional strategy with or without radiation. However, if patients with a low Lung-molGPA (0–1),²⁶ best supportive care is probably the better treatment.

It is crucial to derive benefit from an aggressive treatment approach for patients with BM. While validating this model, we found that OS and PFS of ADC patients with symptomatic BM were significantly affected. The MST in patients scoring 2.5–3 and 1.5–2 with asymptomatic BM

was 28 months and 21 months, respectively, while OS was 39 months and 32.5 months in patients with symptomatic BM. Hence, the treatment of patients with asymptomatic BM may be overlooked. The predictive ability of Lung-molGPA can be considered an estimate of the expected survival on an individual level, which aids the physician in providing an accurate estimation of prognosis to the patient. Meanwhile, it can also be used as a tool to help determine the most suitable treatment and intensity of follow-up.

However, some limitations are inevitable as this was a retrospective study. First, the number of patients with non-ADC was small. Second, few patients underwent genetic testing. Nevertheless, the survival between scores 0–1 and 2.5–3/3.5–4 was statistically significant, which is sufficient to support the feasibility of the clinical application of the Lung-molGPA score. Third, all patients used in this study were from a single institution; thus, the data presented herein might not reflect the reality of other centers. A larger population from multiple institution is needed to balance the variations among various hospitals to further validate the value of Lung-molGPA.

Conclusion

In conclusion, the Lung-molGPA was verified in our retrospective study and was demonstrated to be a useful clinical tool in clinical decision making regarding patient treatment and predict prognosis. However, with the development of novel checkpoint inhibitors and related predictive indicators, static models are no longer suitable. Hence, this model is still exploratory and needs further external validation. In the future, combining more potential indicators, such as PD-L1 expression and markers related to immunity, will improve the model and further guide treatment options to better predict prognosis. More samples are needed to detect and verify the effects of BM symptoms in NSCLC patients.

Ethics Approval

This study was approved by the ethical review board of Shandong Cancer Hospital and Institute and complied with the Helsinki declaration. Written informed consent was obtained from all participants.

Author Contributions

Ji Li and Wang Jing made a significant contribution to the work reported, whether that is in the conception, study design, execution, acquisition of data, analysis and interpretation, or in all these areas. Xiaoyang Zhai and

Wenxiao Jia have drafted or written, or substantially revised or critically reviewed the article. Hui Zhu have agreed on the journal to which the article will be submitted. Hui Zhu and Jinming Yu reviewed and agreed on all versions of the article before submission, during revision, the final version accepted for publication, and any significant changes introduced at the proofing stage. Jinming Yu agree to take responsibility and be accountable for the contents of the article. All authors contributed to data analysis, drafting or revising the article, have agreed on the journal to which the article will be submitted, gave final approval of the version to be published, and agree to be accountable for all aspects of the work.

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Disclosure

The authors report no conflicts of interest in this work.

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