

Sarcopenia and high NLR are associated with the development of hyperprogressive disease after second-line pembrolizumab in patients with non-small-cell lung cancer

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Abstract

The aim of this multicenter retrospective study was to evaluate the incidence of hyperprogressive disease after treatment with pembrolizumab as a second-line treatment in patients (n=167) with non-small-cell lung cancer (NSCLC) with metastatic disease whose tumors expressed programmed death-ligand-1 in [?]1% and to search for factors associated with its development. All patients received platinum-containing chemotherapy as a first-line treatment. The neutrophil to lymphocyte ratio (NLR), platelet to lymphocyte ratio (PLR), and their derivations were retrospectively analyzed. The psoas major muscle area (PMMA) was calculated at the L3 position on computed tomography before chemotherapy and immunotherapy. Patients with [?]PMMA[?]10% were considered to have sarcopenia (low muscle mass). We also performed multinomial logistic regression to estimate the effects of hematological biomarkers and [?]PMMA on the response to immunotherapy. Hyperprogressors (HPs) had significantly higher NLRs, PLRs and [?]PMMA levels than the other patients. Moreover, in multivariate regression analysis, higher levels of [?]PMMA were associated with a decreased likelihood of being a progressor (P) (OR, 0.81; 95% CI, 0.65-0.99; p=0.047) or a nonprogressor (NP) (OR, 0.76; 95% CI, 0.62-0.94; p=0.012) vs an HP. In multivariate analysis, higher NLRs tended to decrease the likelihood of being a P vs an HP (OR, 0.66; 95% CI, 0.42-1.06; p=0.09) and significantly decrease the likelihood of being an NP vs an HP (OR, 0.44; 95% CI, 0.28-0.69; p<0.0001). Our data suggest that a high pre-immunotherapy NLR and the presence of sarcopenia are potential risk factors for the development of hyperprogressive disease.

Introduction

Lung cancer is the leading cause of cancer death worldwide [1], with non-small-cell lung cancer (NSCLC) accounting for approximately 85% of lung cancers. Recently, immunotherapy has represented a breakthrough in oncology, especially in its promise to treat a broad range of advanced cancer types, including NSCLC [2].

Human immune checkpoint inhibitor antibodies inhibit the programmed death (PD-1) receptor or its ligand PD-L1 and thus restore an efficient antitumour T cell response.

Despite advances in the therapeutic landscape of advanced NSCLC without targetable oncogenic driver alterations regarding immunotherapy, the indication spectrum of these new treatments as monotherapy still includes a minority of patients, whereas the vast majority are inevitably candidates for chemotherapy [3, 4]. In the phase II/III KEYNOTE-010 study, pembrolizumab significantly prolonged overall survival (OS) over docetaxel as second-line therapy in advanced NSCLC [5]. Despite these advances in treatment and the increased knowledge of the molecular pathways, there are still challenges in the identification of those patients who are most likely to benefit and those who will not [6, 7]. Durable responses can be observed in some populations, although the percentage has often been found to be approximately 20% [8, 9].

Overexpression of PD-L1 is an important and widely explored predictive biomarker for the response to anti-PD-1/PD-L1 antibodies [10]. Previous studies have demonstrated that the tumor microenvironment, with its most important players being neutrophils, platelets, macrophages and regulatory T cells, plays an essential regulatory role in cancer progression, metastasis and outcome [11, 12]. The results from recent studies suggest that a high neutrophil to lymphocyte ratio (NLR) and platelet to lymphocyte ratio (PLR) may predict a poor response to immune checkpoint inhibitors (ICIs) and poor outcome in patients with NSCLC [13-15]. A recent study proposed that the development of sarcopenia (low muscle mass), measured by the change in the psoas major muscle area (PMMA) at the L3 position, is a negative indicator for the ICI response [16]. In addition to poor responses, immunotherapy was also associated with rapid disease progression, i.e., hyperprogressive disease (HPD), in subpopulations of patients [17] with different incidences [18]. Unfortunately, currently, there are no biomarkers that predict the development of this life-threatening condition.

The purpose of this retrospective study was to evaluate the incidence of HPD after treatment with pembrolizumab as a second-line treatment in metastatic NSCLC patients and to search for indicators that are associated with the development of HPD.

Materials and Methods

Patient selection

In this retrospective cohort study, we reviewed the cases of 167 patients from five centers in Bulgaria with metastatic NSCLC treated with pembrolizumab after progression upon first-line platinum-based chemotherapy between April 2017 and February 2020. The procedure was approved by the Scientific Research Ethics Committee at the Hospital “Nadezhda” in Sofia. The eligibility criteria were as follows: (1) age \geq 18 years old, (2) histologically confirmed diagnosis of NSCLC in the metastatic stage, (3) wild-type epidermal growth factor receptor/anaplastic lymphoma kinase, (4) Eastern Cooperative Oncology Group-Performance status (ECOG-PS) $<$ 2, (5) disease progression after receiving one prior platinum-based systemic therapy for metastatic disease with measurable disease lesions, (6) available blood cell count and blood samples, and (7) available computed tomography (CT) scans. Immunotherapy were administered after at least 3 weeks after previous treatment. Patients were excluded if they had brain metastases (since corticosteroid use may compromise therapy), autoimmune disease, symptomatic interstitial lung disease, systemic immunosuppression, or prior treatment with immune-stimulatory antitumor agents, including checkpoint inhibitors. Patients did not show any clinical or computed tomography signs of active infection. Tumor PD-L1 status was assessed. Pembrolizumab was initially administered via 2 mg/kg intravenous (i.v.) injection over 60 minutes every 3 weeks and later via 200 mg i.v. injection (flat dose) every 3 weeks.

Data collection

Data collected included demographics, PD-L1 status, metastatic sites, description of first-line treatment, date of progression as determined by radiology reports, and date of death or last follow-up. Peripheral blood

samples were collected from patients included in the study the day of first-line chemotherapy administration at baseline and the day of first immunotherapy infusion upon progression. Of interest were the following hematological and biochemistry parameters: absolute neutrophil count (ANC), absolute lymphocyte count (ALC) and platelet count (APC), which enable calculation of the NLR (neutrophil to lymphocyte ratio: ANC/ALC) and the PLR (platelet to lymphocyte ratio: APC/ALC). NLR1 and PLR1 were calculated before the first cycle of chemotherapy, and NLR2 and PLR2 were calculated before the first pembrolizumab infusion. Δ NLR (NLR2-NLR1) and Δ PLR (PLR2-PLR1) were calculated. An NLR>5 was considered high in accordance with earlier reports [19-21]. The median value of NLR2 was 4.8. The median value (174) of the PLR was used to group cases into two categories of low (\leq median) and high ($>$ median) PLRs.

The tumor PD-L1 protein expression was analyzed by immunohistochemistry of tumor cells in archived biopsy samples of tumors, and the cutoff for positivity was 1%. In addition, we subdivided the positive group into expression categories: expression in 50%, expression in 25-49%, and expression in 1-24%.

Measurement of psoas major muscle area

The psoas major muscle area was calculated at the L3 position on computed tomography. The PMMA was calculated before chemotherapy and before pembrolizumab infusion. We were able to measure Δ PMMA in only 112 patients of the whole cohort, since the rest were staged only with a CT scan of the thorax and upper abdomen and the area of the patient's psoas major muscle at the L3 position was not available. The PMMA was measured in the region of interest by tracing an outline using the image viewer software "DICOM". The following formula was used: % change of PMMA = $(\frac{1-PMMA \text{ before P}}{PMMA \text{ before CT}}) * 100$. Patients with a change in PMMA $\geq 10\%$ were considered to have sarcopenia [16].

Endpoints

The tumor response was assessed according to the Response Evaluation Criteria in Solid Tumors (ver. 1.1) (RECIST 1.1), and clinical tumor response was assessed every 3 months or at clinical deterioration. Hyperprogression was defined if at least three of the following existed: 1. time to treatment failure < 3 months; 2. increase $\geq 50\%$ in the sum of target lesion major diameters between baseline and first radiological evaluation; 3. appearance of at least two new lesions in an organ already involved between baseline and first radiological evaluation; 4. spread of the disease to a new organ between baseline and first radiological evaluation; and 5. clinical deterioration with a decrease in ECOG PS ≥ 2 during the first three months of treatment. Pseudoprogression was defined as initial progression followed either by partial response or stable disease lasting at least 6 months. OS was defined as the interval between diagnosis of the disease and death or the date of the last follow-up evaluation.

Statistical design and analysis

Data were managed and analyzed using SPSS software ver. 23. The demographic characteristics were expressed as frequencies and percentages for categorical variables and as medians and means with standard deviations for quantitative variables. The Mann-Whitney U test, Spearman correlation and χ^2 test were used to compare and evaluate the correlations between the biomarkers and the clinicopathological characteristics of the patients, such as age, sex, the NLR, and the PLR. To assess the correlations between test results, rho values were interpreted as follows: < 0.39 , weak correlation; 0.40-0.59, moderate correlation; 0.60-0.79, strong correlation; and ≥ 0.80 , very strong correlation. The Kruskal-Wallis one-way analysis of variance was used to compare the levels of hematological biomarkers, Δ PMMA and response to pembrolizumab at the first CT scan. The Wilcoxon and McNemar tests were used to compare quantitative and categorical biomarker values and their derivations. The diagnostic accuracy of biomarkers was determined by obtaining the largest possible area under the curve (AUC) in receiver operating characteristic curve (ROC) analysis. AUC values ≥ 0.9 were considered "excellent", ≥ 0.80 were considered "good", ≥ 0.7 were considered "fair" and < 0.70 were considered "poor". Survival curves according to the response on the first CT scan were estimated using the Kaplan-Meier method, and differences were assessed using the log-rank test. We also

performed multinomial logistic regression to estimate the effects of hematological biomarkers and [?]PMMA on the response to treatment. Two-tailed p-values <0.05 were considered significant.

Results

Baseline characteristics

This study included 167 patients who received anti-PD1 treatment with pembrolizumab after failure of first-line chemotherapy. The clinical characteristics of the patients and relations with response on the first CT scan are summarized in Table 1. The mean age was 60.2 ± 6.8 years; most of the patients were men (64%), and almost all patients exhibited an ECOG PS of 1 (98%). The lung was the most common metastatic site (73%), followed by pleural effusion (59%) and bone (40%). All patients were eligible for the examination of tumor PD-L1 expression, of which 13 patients (7.8%) had expression in more than 50%, 80 patients (47.9%) had expression in 25-49%, and 74 patients (44.3%) had expression in 1-24%. Of all the clinicopathological characteristics of the patients, only NLR2, PLR2 and the presence of sarcopenia were significantly related to the response on the first CT scan (Table 1).

Hematological biomarkers and their relation to response on the first CT scan

On the first CT scan after chemotherapy treatment, 15 (8.9%) patients showed progressive disease. After treatment with pembrolizumab on the first CT scan evaluation, these 15 patients were subdivided as follows: 8 hyperprogressors (HPs), 1 pseudoprogessor (PP) and 6 nonprogressors (NPs). These 15 patients had significantly higher NLR1 and PLR1 than the patients without progressive disease (7.49 ± 2.8 vs 4.31 ± 2.45 ; 283.3 ± 96.5 vs 207 ± 102.6 , respectively). Twelve of them had an $NLR > 5$, and at least 9 ([?]PMMA was not available for 3) of them had [?]PMMA [?]10%.

On the first CT scan after immunotherapy treatment, 45 (26.9%) patients showed progressive disease, and at least 25 ([?]PMMA was not available for 7) of them had [?]PMMA [?]10%. Of them, 16 patients (9.6%) were classified as HPs, 5 (2.9%) were classified as PPs, and the remaining 24 (14%) were classified as Ps. Patients with pseudoprogression were without any clinical deterioration and received further treatment with immunotherapy for another 8 weeks, when the control CT scan proved a partial response for 3 patients and stable disease for 2 patients; the response lasted for at least 6 months. Of all HPs, 15 (93%) had an $NLR > 5$. HPs had higher mean values of NLR2, PLR2 and [?]NLR, but not higher [?]PLR values, than Ps or NPs (Table 2). There was no significant difference in hematological parameters between HPs and PPs, Ps and NPs, Ps and PPs, or NPs and PPs, except for NLR2, for which NPs had significantly lower values than PPs (Table 2).

ROC analysis was performed to explore the potential predictive role of these biomarkers, NLR2, PLR2, ΔNLR , and ΔPLR , as noninvasive biomarkers for discrimination between patients with or without HPD (Table 3). At the optimal cutoff values for NLR2, the biomarker could significantly and well distinguish between patients with or without HPD (AUC = 0.85, 95% CI: 0.75- 0.95, $p < 0.001$) with a sensitivity of 87.5% and a specificity of 68.9%. PLR2 also allowed significant but fair discrimination between patients with and without HPD (AUC = 0.79, 95% CI: 0.66- 0.92, $p < 0.001$) with a sensitivity of 75.0% and a specificity of 64.1% (Figure 1A and B). ΔNLR could also discriminate between patients with and without HPD, but poorly (Table 3). The Wilcoxon test showed that the ALC and APC did not change significantly from chemotherapy. Nevertheless, the ANC significantly differed between the first cycle of chemotherapy and the first pembrolizumab infusion. The McNemar test showed that the proportion of patients with an $NLR > 5$ and a high PLR did not change significantly with chemotherapy treatment.

A significantly strong correlation was detected between NLR1 and PLR1 ($\rho=0.763$) and NLR2 and PLR2 ($\rho=0.785$), and a moderate correlation was detected between ΔNLR and ΔPLR ($\rho=0.465$).

Sarcopenia and its relation to response on the first CT scan

The number of patients who developed sarcopenia during chemotherapy treatment was 34 (30.3%). After treatment with pembrolizumab after the first CT scan evaluation, these patients were subdivided as follows: 15 were HPs, 4 were PPs, 7 were Ps, and 8 were NPs. There was a significant relationship between the presence of sarcopenia and the response on the first CT scan (Table 1).

The Kruskal-Wallis one-way analysis of variance showed that there were significant differences in [?]PMMA only between HPs and Ps (16.2 ± 4.8 vs 8.3 ± 8.1 ; $p=0.009$) and NPs and HPs (5.8 ± 13.8 vs 16.2 ± 4.8 ; $p<0.0001$). The Mann-Whitney U test showed that patients with sarcopenia had significantly higher NLR2 and PLR2 values than patients without sarcopenia (7.9 ± 3.2 vs 3.6 ± 2.3 and 315.9 ± 157.9 vs 168.7 ± 93.8 , respectively; both $p<0.0001$). ROC analysis revealed that at the cutoff value of Δ PMMA [?]10% could distinguish between patients with or without HPD with an AUC=0.89 (95% CI, 0.82-0.96; $p<0.0001$), and with a sensitivity of 93.8% and specificity of 79.2% (Figure 1C). After adjustment for age, sex, PD-L1 expression, number of metastatic sites, NLR, PLR and their derivations, higher levels of [?]PMMA were associated with a decreased likelihood of being a P vs an HP (OR, 0.81; 95% CI, 0.65-0.99; $p=0.047$) and being an NP vs an HP (OR, 0.76; 95% CI, 0.62-0.94; $p=0.012$).

A significant but weak correlation was detected between [?]PMMA and NLR2 ($\rho=0.365$), [?]PMMA and PLR2 ($\rho=0.279$), and [?]PMMA and age ($\rho=0.292$).

Association between hematological biomarkers and response on the first CT scan

In univariate analysis, higher levels of NLR2 and PLR2 were associated with a decreased likelihood of being a P vs an HP (OR, 0.67; 95% CI, 0.53-0.86; $p=0.001$; OR, 0.993; 95% CI, 0.98-0.99; $p=0.006$, respectively). On univariate analysis, higher levels of NLR2, PLR2 and [?]PLR were associated with a decreased likelihood of being an NP vs an HP (OR, 0.61; 95% CI, 0.48-0.76; $p<0.0001$; OR, 0.991; 95% CI, 0.98-0.99; $p<0.0001$, OR, 0.994; 95% CI, 0.98-0.99; $p=0.008$, respectively). After adjustment for age, sex, PD-L1 expression and number of metastatic sites, higher levels of NLR2 were associated with a decreased likelihood of being an NP vs an HP (OR, 0.44; 95% CI, 0.28-0.69; $p<0.0001$) and showed a trend for being associated with a decreased likelihood of being a P vs an HP (OR, 0.66; 95% CI, 0.42-1.06; $p=0.09$) (Table 4).

Overall survival of HPs, PPs, Ps, NPs and patients with sarcopenia

HPs at the first CT evaluation had a significantly shorter mean OS (9.83 months; 95% CI 8.44-11.22) than PPs (19.18 months; 95% CI 14.13-24.22) (log-rank test $p=0.001$), Ps (17.32 months; 95% CI 15.67-18.98) (log-rank test $p<0.001$) and NPs (29.79 months; 95% CI 26.87-32.71) (log-rank test $p<0.001$) (Figure 2A). Patients with sarcopenia had a significantly shorter mean OS (13.5 months; 95% CI 11.7-15.2) than patients without sarcopenia (31.5 months; 95% CI 27.6-35.8) (log-rank test $p<0.001$) (Figure 2B).

Discussion

The current study found that the incidence of HPD after treatment with pembrolizumab as a second-line therapy was 9.6%. Higher NLRs and [?]PMMA before treatment with pembrolizumab were associated with a higher risk for the development of HPD. Our results suggest for the first time that in the population of patients who do not respond early to platinum-based chemotherapy, some patients who are at risk for the development of HPD on immunotherapy exist.

ICIs may promote tumor growth kinetics in certain patients and lead to the development of HPD with an incidence in the range from 4% to 29% in different studies [17, 22] There is no uniform definition for HPD, even though some authors still doubt its existence because it is an ad hoc observation [23]. The majority of researchers rely on the rate of target lesion growth for defining HPD [17, 24, 25]. Since this underestimates the rate of development of metastasis, we included in our study involvement of new lesions as a part of the

definition for HPD. In conjunction with others, our results suggest that a dramatic increase in tumor growth induced by ICIs is restricted to a relatively small subpopulation of patients, approximately 10% [17, 22]. We could not find any association between HPD and age, sex, PS, PD-L1 status, or number of metastatic sites.

Although how and why HPD occurs are still unclear, our results shed some light on this problem. Our results show that the NLR is an important risk factor for the development of HPD. It is well known that high baseline NLR and PLR values and their derivations are linked significantly with worse OS and PFS in patients with NSCLC treated with ICIs [6, 15]. Although there is no clear explanation for this phenomenon, neutrophils and platelets may promote tumor progression as well as metastases by exercising a direct effect on tumor cells or by indirectly affecting other components of the tumor microenvironment [26, 27]. This effect is achieved through the secretion and release of various chemokines and cytokines, including transforming growth factor-beta, vascular endothelial growth factor, IL-6, IL-8, matrix metalloproteinases and the formation of neutrophil extracellular traps [27]. Neutrophils are considered the most important inflammatory cell population in the tumors of NSCLC patients and promote metastasis, thus potentially compromising the antitumor immune response [28]. Recent studies reported that blood neutrophils, identified by the NLR, were directly linked with the number of intratumoral neutrophil populations, which may have the potential to compromise the antitumor immune response [29]. Lower counts of lymphocytes usually reflect an impairment of cell-mediated immunity. It has been shown that increased infiltration of lymphocytes in the tumor region is associated with better responsiveness to treatment and prognosis in patients with solid tumors [30]. To further complicate the results, it has been shown that activation of tumor lymphocytes could trigger local inflammation and matrix and metabolism modifications that could lead to tumor escape [31]. Moreover, in murine models, it has been shown that HPD is associated with vast infiltration in the tumor microenvironment of primary myeloid cells (mostly precursors of neutrophils and macrophages), which express high levels of activation and inhibitory receptors [32].

Consistent with others, our research finds a positive relationship between sarcopenia and the NLR [33, 34]. Cancer-associated cachexia is a well-known negative prognostic marker with an incidence of up to 40% in the cancer population [35]. Although there are differences in the definitions used for cachexia and sarcopenia, they are often indistinguishable in clinical practice, and cachexia and its key feature inflammation could lead to sarcopenia [36]. It was shown that sarcopenia measured via [?]PMMA with a CT scan at the L3 position is much more reliable than body mass index, and it is widely used, with a variety of cutoff values [16, 37, 38]. A recent report showed that patients receiving immunotherapy may be particularly susceptible to cancer-associated cachexia [39]. This may explain at least in part why in some patients with pre-existing sarcopenia, immunotherapy may induce rapid tumor growth; upregulation of stress hormone production together with preexisting systemic immunosuppression and the presence of high inflammatory markers such as NLR accelerate tumor growth and thus ultimately lead to HPD.

Several limitations were identified in our study. First, our study is retrospective and has a relatively small sample size; therefore, there is potential for bias. Moreover, the predictive value of the NLR and sarcopenia was not compared to that of some genetic predictive markers, such as *EGFR*, *MDM2* and *DNMT3A*. Similarly, tumor mutational burden was not available. Second, in our study, only 112 patients (67%) were available for evaluation of the [?]PMMA. Finally, due to the vague and unclear definitions of HPD in the literature, our definition for this phenomenon might be criticized.

Despite these limitations, our results suggest for the first time that patients with a high NLR and sarcopenia before immunotherapy are at higher risk for hyperprogression and short overall survival. This may be helpful to clinicians in their choice of treatment, especially for patients who progressed rapidly on platinum-based chemotherapy and with high NLRs and sarcopenia; perhaps a combination of chemotherapy and immunotherapy or new molecules in clinical trials could be used for these patients. Drugs that are capable of transforming neutrophils into a functional state with antitumor activity are needed to improve patient outcomes.

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Table 1. Relationship between baseline clinicopathological characteristics of patients and response on the first computed tomography scan - hyperprogressors (HPs), progressors (Ps), and nonprogressors (NPs).

	HPs	Ps	NPs	p value
Sex				
Male	8 (7.5%)	16 (15%)	83 (77.6%)	0.35
Female	8 (14.5%)	8 (14.5%)	39 (70.9%)	
ECOG PS				0.70
0	0 (0%)	1 (25%)	3 (75%)	
1	16 (10.1%)	23 (14.6%)	119 (75.3%)	
Lung metastasis				0.97
No	4 (10.8%)	5 (13.5%)	28 (77.7%)	
Yes	12 (9.8%)	18 (14.8%)	92 (75.4%)	
Liver metastasis				0.35
No	10 (8.4%)	16 (13.4%)	93 (78.2%)	
Yes	6 (15%)	7 (17.5%)	27 (67.5%)	
Pleural effusion				0.85
No	5 (8.3%)	9 (15%)	46 (76.7%)	
Yes	11 (11.1%)	14 (14.1%)	74(74.7%)	
Bone metastasis				0.92
No	10 (10.9%)	13 (14.1%)	69 (75%)	
Yes	6 (9%)	10 (14.9%)	51 (76.1)	
Number of metastatic lesions				0.45
<2	7 (7.5%)	14 (15.1%)	72 (77.4%)	?;?
2	9 (13.8%)	9 (13.8%)	48 (72.7%)	
PD-L1 expression				0.71
1-24%	8 (11.6%)	12 (17.4%)	49 (71%)	
25-49%	6 (7.5%)	10 (12.5%)	64 (80%)	
>50%	2 (15.4%)	2 (15.4%)	9 (69.2%)	
NLR				<0.0001?;?
5	1 (1.2%)	10 (11.8%)	74 (87.1%)	
>5	15 (19.5%)	14 (18.2%)	48 (62.3%)	
PLR				0.004?;?
median	3 (3.6%)	9 (10.8%)	71 (85.5%)	
>median	13 (16.5%)	15 (19%)	51 (64.6%)	
Age				0.71
<65 years	9 (8.5%)	16 (15.1%)	81 (76.4%)	?;?
65 years	7 (12.5%)	8 (14.3%)	41 (73.2%)	
Sarcopenia				<0.0001
No	1 (1.3%)	11 (14.3%)	65 (84.4%)	
Yes	15 (50%)	7 (23.3%)	8 (26.7%)	

Table 2 . Comparison between neutrophil to lymphocyte ratio (NLR), platelet-lymphocyte ratio (PLR), their derivations and response on the first computed tomography scan- hyperprogressors (HPs), pseudoprogressors (PPs), progressors (Ps), and nonprogressors (NPs). Means with standard deviations are shown. Adjusted p-values were used.

	NLR2	p	PLR2	p	[?]NLR	p	[?]PLR	p
HPs vs Ps	9.01±2.66	0.006	368.29±193.8	0.044	1.29±3.02	0.04	65.67±185.31	0.44
vs			vs		vs 0.09±2.3		vs	
	5.31±3.26		218.53±108.01				24.15±99.82	

	NLR2	p	PLR2	p	[?]NLR	p	[?]PLR	p
HPs vs PPs	9.01±2.66	0.91	368.29±193.8	0.72	1.29±3.02	0.37	65.67±185.31	0.78
	vs		vs		vs		vs	
	8.39±1.32		303.09±115.69		2.28±1.5		17.53±59.75	
HPs vs NPs	9.01±2.66	<0.0001	368.29±193.8	<0.0001	1.29±3.02	0.017	65.67±185.31	<0.0001
	vs		vs		vs		vs	
	4.35±3.01		192.67±109.34		0.45±2.01		-9.99±85.88	
Ps vs NPs	5.31±3.26	0.29	218.53±108.01	0.36	0.09±2.3	vs 0.28	24.15±99.82	0.36
	vs		vs		0.45±2.01		vs	
	4.35±3.01		192.67±109.34				-9.99±85.88	
Ps vs PPs	5.31±3.26	0.25	218.53±108.01	0.97	0.09±2.3	vs 0.041	24.15±99.82	0.98
	vs		vs		2.28±1.5		vs	
	8.39±1.32		303.09±115.69				17.53±59.75	
NPs vs PPs	4.35±3.01	0.040	192.67±109.34	0.31	0.45±2.01	0.10	-9.99±85.88	0.30
	vs		vs		vs 2.28±1.5		vs	
	8.39±1.32		303.09±115.69				17.53±59.75	

Table 3. Receiver operating curve (ROC) curve analysis was performed using the neutrophil to lymphocyte ratio (NLR), platelet-lymphocyte ratio (PLR) and their derivations to differentiate between patients with and without hyperprogressive disease. The diagnostic accuracy of biomarkers was determined by obtaining the largest possible area under the curve (AUC) in ROC analysis.

Biomarker	AUC 95% CI	p value	Sensitivity (%)	Specificity (%)
NLR2	0.85 (0.75-0.95)	<0.001	87.5	68.9
PLR2	0.79 (0.66-0.92)	<0.001	75.0	64.1
ΔNLR	0.68 (0.54-0.83)	0.016	62.5	61.3
ΔPLR	0.62 (0.44-0.79)	0.13	56.3	60.7

Table 4. Associations between hematological biomarkers and response to pembrolizumab on the first computed tomography scan- hyperprogressors (HPs), progressors (Ps), and nonprogressors (NPs).

Multinomial outcome	Predictor	Unadjusted	Unadjusted	Unadjusted	Adjusted*	Adjusted*	Adjusted*
		OR	95% CI	p	OR	95% CI	p
Ps vs HPs	NLR2	0.67	0.53-0.86	0.001	0.66	0.42-1.06	0.09
NPs vs HPs	NLR2	0.61	0.48-0.76	<0.0001	0.44	0.28-0.69	<0.0001
Ps vs HPs	PLR2	0.993	0.98-0.99	0.006	0.992	0.98-1.002	0.12
NPs vs HPs	PLR2	0.991	0.98-0.99	<0.0001	0.999	0.99-1.008	0.82
Ps vs HPs	[?]NLR	0.74	0.54-1.02	0.066	1.10	0.67-1.79	0.69
NPs vs HPs	[?]NLR	0.79	0.61-1.02	0.070	0.69	0.38-1.24	0.22
Ps vs HPs	[?]PLR	0.997	0.99-1.002	0.29	1.005	0.99-1.015	0.36
NPs vs HPs	[?]PLR	0.994	0.98-0.99	0.008	0.994	0.98-1.003	0.16

*Adjusted for age, sex, PD-L1 expression and number of metastatic sites

Figure Legends

Figure 1. Receiver operating curve (ROC) curve analysis using the neutrophil to lymphocyte ratio (NLR), the platelet-lymphocyte ratio (PLR) and [?]PMMA to differentiate between patients with and without hyper-

progressive disease. The diagnostic accuracy of biomarkers was determined by obtaining the largest possible area under the curve (AUC) in ROC analysis **A**. NLR2 AUC=0.85; **B**. PLR2 AUC=0.79; **C**. [?]PMMA AUC=0.89.

Figure 2. Kaplan-Meier estimates of overall survival (OS) in hyperprogressors (HPs), pseudoprogressors (PPs), progressors (Ps), nonprogressors (NPs) and patients with sarcopenia. **A**. HPs had a significantly shorter mean OS (9.83 months; 95% CI 8.44-11.22) than PPs (19.18 months; 95% CI 14.13-24.22) (log-rank test $p=0.001$), Ps (17.32 months; 95% CI 15.67-18.98) (log-rank test $p<0.001$) and NPs (29.79 months; 95% CI 26.87-32.71) (log-rank test $p<0.001$). **B**. Patients with [?]PMMA \geq 10% had a significantly shorter mean OS (13.5 months; 95% CI 11.7-15.2) than patients with [?]PMMA $<$ 10% (31.5 months; 95% CI 27.6-35.8) (log-rank test $p<0.001$).

Figure 1.

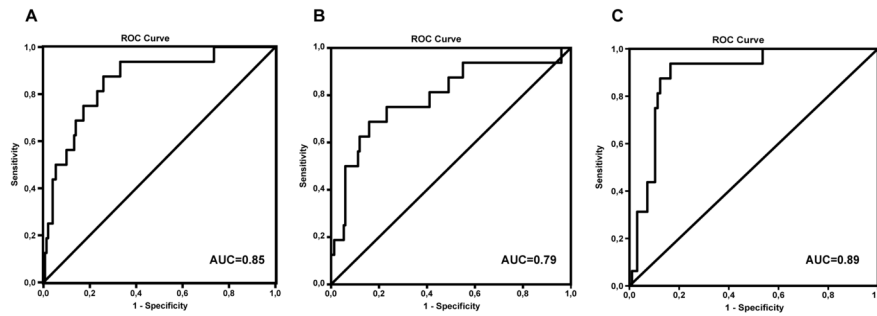


Figure 2.

