

Prognostic value of the neutrophil-to-lymphocyte ratio and primary tumor location in patients with EGFR-mutated metastatic non-small cell lung cancer treated with TKIs

Note: We found that we got a wrong results in the previous article. So, we checked our collected database and calculated again and corrected now.

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Abstract

Background: Targeted therapy with the epidermal growth factor receptor (EGFR) tyrosine kinase inhibitors (TKIs) has improved the field of metastatic non-small cell lung cancer treatment. Higher neutrophil-to-lymphocyte ratio (NLR) and lower relative lymphocyte counts as inflammatory indicators and associated with worse overall survival and progression free survival (PFS) in several tumor types. Few studies focused on these inflammation markers in field of TKIs eras.

Methods: Patients with advanced EGFR mutation NSCLC treated with TKIs were included. Pre-treatment NLR means neutrophil to lymphocyte ratio measured in peripheral blood within one week before treating with TKIs. The baseline clinical characteristics of each group were compared by chi-square and t tests. Cox regression analyses were used to evaluate prognostic value of peripheral blood parameters on progression free survival (PFS). All prognostic factors were explored with multivariable regression.

Results: We retrospectively analyzed 221 patients with metastatic NSCLC harboring exon 19 deletion, 21 L858R or rare mutation and receiving TKIs. Finally, a total of 190 patients were analyzed. The optimal cutoff values for pretreatment absolute lymphocyte count (Lym), lymphocyte percentage (Lym%), absolute neutrophil count (Neu), the percentage of neutrophil granulocytes (Neu%) and NLR were 1.625, 18.8%, 3.675, 51.8% and 4.965, respectively. Patients with high neutrophil percent (18.8 months vs 13.0 months, $P=0.003$), absolute neutrophil counts (12.0 months vs 14.5 months, $P=0.014$) and NLR (7.0 months vs 15.2 months, $P<0.001$, 1-year PFS Rate, 38.4%, respectively) had worse PFS. In contrast, patients with high absolute lymphocyte counts (16.5 months vs 13.0 months, $P=0.012$) and lymphocyte percent (15.3 months vs 8.8 months, $P<0.001$) had a better progression-free-survival. Besides, tumor location is also an important factor for prognosis (14.3 months vs 11.6 months, $P=0.003$). On multivariate analysis, NLR and primary tumor location were both identified as independent and significantly risk indicators for worse PFS.

Conclusion: NLR and primary location are both independent prognostic factors for PFS in patients with metastatic EGFR mutated lung tumor. Whether or not NLR and primary location could be useful markers in efficacy prediction of TKIs in advanced NSCLC calls for further investigation. **KEYWORDS:** Epidermal growth factor; NSCLC; TKI; Primary location; NLR; PFS

1. Background

Lung cancer continues to be the leading cause of cancer-related deaths worldwide, especially in patients with metastatic disease [1]. Non-small cell lung cancer (NSCLC) as the most common type of lung carcinoma, accounts for approximately 85%-90% of all lung cancer cases. Lung neoplasms can also be grouped into the central and peripheral subtypes according to primary tumor location [2]. Pulmonary adenocarcinoma pathological subtypes have replaced squamous cell carcinoma in recent years, often located in the periphery, as one of the most common cell types of lung cancer. In recent years, a number

of previous studies have shown that primary tumor location is an important factor to guide treatment schedules and predict clinical prognosis in lung tumors[3-6].

Advances in epidermal growth factor receptor tyrosine kinase inhibitors EGFR-TKIs of non-small cell lung cancer (NSCLC) have led to a new era of target therapy, particularly in patients with NSCLC who have EGFR mutations. Until recently, targeted therapy has remained the first-line treatment for the majority of patients with targetable oncogenic driver alterations. In recent years, gefitinib, erlotinib, afatinib, and osimertinib have shown better clinical outcomes and response rates than chemotherapy using cytotoxic drugs.

Host immunity may affect prognosis in patients with various cancers. Evidence has demonstrated that immune system plays an important role in promoting anti-tumor defense. Tumor-associated inflammation and tumor microenvironment play a critical role in cancer development, progression and metastasis[7-9]. The neutrophil-to-lymphocyte ratio (NLR) is one of the most widely used inflammation biomarkers in solid cancers and can be easily calculated from routine blood examination results. Inflammation not only can contribute to development of various cancers, but also is recognized as a hallmark of cancer[10]. For example, several peripheral blood parameters, including markers of systemic inflammation such as baseline the NLR, the lymphocyte-to-monocyte ratio (LMR) and absolute lymphocyte count (Lym) have been associated with survival in patients with metastatic lung neoplasms treated with the ICIs[11, 12].

Neutrophil counts, lymphocyte counts and NLR play significant roles in the inflammatory response and demonstrates their value in various of solid tumors. As people costs much in treatments, inexpensive, readily available and more effective potential prognostic markers need to be used to assist the prognosis and risk stratification of patients with the lung cancer. Previous studies have shown that the NLR can be considered as a predictor to evaluate the prognosis, which can be used in the EGFR-TKIs and ICIs [13-15]. Few studies have found an association between NLR and OS. There is a lack of understanding of the relationship between peripheral blood counts and progression-free survival (PFS), particular in patients treated with TKIs.

The aim of our study was to find the prognostic value of pre-treatment complete blood count (CBC) parameters in NSCLC patients with advanced NSCLC treated with EGFR-TKIs (as first-line to third-line therapy). Additionally, we needed to find various determined factors to predict clinical outcomes. In general, the main objective was to explore more effective, useful and noninvasive predictors to assess the benefits of patients receiving TKI treatment.

2. Methods

2.1 Patients and clinical characteristics

This retrospective study was approved by review board of Shandong Cancer Hospital and Institute. We respectively analyzed patients with metastatic or recurrent postoperative NSCLC. All patients needed to

meet the following standard criteria: 1) were of 18 years or older, 2) pathologically proven adenocarcinoma with a positive EGFR mutation test before starting any treatment, including chemotherapy, surgery, radiotherapy (RT) and targeted therapy, 3) complete medical records/computed tomography (CT) scans of the chest and/or positron emission tomography (PET) scans/bronchoscopies, 4) treatment with TKI drugs as the first-, second- or third-line therapy, 5) results of complete peripheral blood test, including neutrophil and lymphocyte counts, within 1 week before receiving EGFR-TKI treatment and 6) received TKI drugs including Gefitinib, Erlotinib and Icotinib. Thus, patients who meet all the above-mentioned criteria were included from the electronic record system.

The clinical stage was determined by the 7th edition of the American Joint Committee on Cancer (AJCC)/Union for International Cancer Control (UICC) staging system. Pre-treatment NLR means the nearest peripheral absolute neutrophil and absolute lymphocyte ratio before starting of TKIs within 1 week. PFS was defined as the time from treatment with the EGFR-TKIs to the terminology or pathology evidence of progression or recurrence. OS was measured from the day of NSCLC diagnosis to the date of death from any.

2.2 EGFR mutation test

A total of 190 specimens before receiving EGFR-TKIs were obtained via tissue biopsy, including bronchoscopy, CT-guided biopsy, or surgical procedures from primary or metastatic sites. If we could not obtain the tissues, we collected peripheral blood and metastatic body fluid. The majority of samples use the peptide nucleic acid (PNA)-mediated polymerase chain reaction (PCR) clamping method. Few samples were sequenced with targeted next-generation sequencing (NGS) of 18 lung cancer gene panels.

2.3 Tumor location evaluation

According to results of patients' CT imaging, bronchoscopy, or both, we identified tumor as central and peripheral lung cancer. As there is no standard definition to classify peripheral and central lung tumor. We defined central tumors as occurring from segmental or proximal bronchi. As for peripheral type, we considered that tumors in subsegmental or other distal bronchi and bronchioli according to previous studies¹². All images were anonymized and blindly evaluated by one radiologist and one oncologist. For discordant cases, a second oncologist would evaluate the images.

3. Statistical Analyses

Pre-treatment blood data were obtained from electronic medical records. Student's t-test was used to evaluate the difference in absolute lymphocyte counts, relative lymphocyte counts and relative neutrophil counts in two different groups. We used Mann-Whitney U test for the ordinal data and Pearson's χ^2 or Fisher's exact test to compare clinical characteristics at baseline. EGFR mutation status and primary tumor location (central vs. peripheral) were analyzed using Pearson's chi-square tests. Receiver operating characteristic (ROC) curve analysis was used to determine the optimal value of Lym, percentage of lymphocytes (Lym%), absolute neutrophil count (Neu), the percentage of neutrophil granulocytes (Neu%)

and the NLR in terms of their association with PFS. Patients and clinical characteristics were summarized with descriptive statistics. We also used the Kaplan-Meier method and log-rank for univariate survival analysis. Multivariate Cox regression analysis was used to analyze the effect of different clinical characteristics on PFS and OS, and statistical analyses were performed with SPSS 19.0. $P < 0.05$ was considered statistically significant.

4. Results

4.1 Patients and tumor characteristics

From January 2014 to November 2018, a total of 221 patients were treated with TKIs. Thirty-one patients did not meet the eligible requirements (complete peripheral blood tests were missing for 29 patients, the primary tumour location was difficult to define in the imaging data for 1 patient, and imaging data before starting TKIs were missing for 1 patient). Baseline characteristics of the patients are summarized in Table 1. The majority of patients were female ($N=122$), and the median age was 58.0 years (range: 30 - 87 years), a total of 148 patients had never smoked. A total of 172 patients underwent PCR tests and 18 underwent NGS tests. We also found the most oncogenic alteration was EGFR L858R mutation ($N=110$). Additionally, 83 patients had bone metastasis (43.7%), 26 (13.7%) had liver metastasis, and 59 (31.1%) had brain metastasis. In addition, many patients received chemotherapy as the first-line therapy ($N=90$). 27 patients underwent surgery before targeted therapies. No correlations were found between primary location and EGFR mutation status. (Pearson's chi-square= 0.76, $P=0.963$). 142 patients had peripheral-type adenocarcinoma. There were 164 patients with stage IV lung cancer. A total of 26 patients had postoperative recurrence.

4.2 Optimal cutoff value for CBC

Median PFS was 12.51 months when we used PFS as an end point for blood routine. According to ROC analysis, we obtained the following optimal cut-off values: 1.625 for pre-treatment Lym, 18.8% for pre-treatment Lym%, 3.675 for pre-treatment Neu, 51.8% for pre-treatment Neu% and 4.965 for pre-treatment NLR for PFS (Table 3). Among the many routine peripheral blood indicators, pre-treatment Lym had the largest AUC, which was 63.2%. However, the pre-treatment Neu% had the lowest AUC, which was 49.9%, indicating its low predictive value.

4.3 Association between pre-treatment CBC and clinicopathological factors

As Table 4 demonstrates, a positive association between the dichotomized NLR and Neu was found, with a kappa value of 0.487 ($P < 0.0001$). Additionally, in Table 4, a negative correlation between the dichotomized NLR and Lym were consistently explored, with a kappa value of 0.310 ($P < 0.0001$). In addition, we also found that there were positive correlations between bone metastasis and pre-treatment Neu% and NLR ($P=0.007$ and $P=0.022$, respectively). There were negative correlations between pre-treatment Lym% and bone metastasis ($P=0.029$). The absolute pre-treatment Lym tended to correlate with

PFS (P=0.007). Additionally, there were no correlations between other blood biomarkers and metastasis sites. We did not find a relationship between peripheral blood biomarkers and primary tumour location.

4.4 Prognostic factors for PFS

The median PFS was significantly related to the pre-treatment Lym, Lym%, Neu, Neu% and NLR. In summary, the PFS of patients with a higher Lym (HLym) and higher relative lymphocyte count HLym% was significantly higher than those of the patients with LLym and LLym% (P<0.05; Fig. 1A, B). In contrast, patients with HNeu, HNeu% and HNLR had poorer PFS than those with LNeu, LNeu% and LNLR (P<0.05; Figs. 1C, 1D; and 2). We also found that peripheral-type lung neoplasms had a better clinical outcome than central-type lung neoplasms (P=0.003, hazard ratio (HR)= 1.739, 95% confidence interval (CI)= 1.207-2.506) (Fig. 2). Pre-treatment blood tests were found to be important for PFS, patients with Lym <1.625 had shorter PFS than those with Lym \geq 1.625 (13.0 months and 16.5 months, P=0.012, HR=0.662, 95% CI= 0.480-0.915). Median PFS was 8.8 months in patients with Lym%<18.8%, while the PFS was 15.3 months in patients with

Lym% \geq 18.8%. (P=0.001, HR=0.391, 95%CI=0.274-0.560) Besides, patients with a HNeu, relative neutrophil count and HNLR had a worse clinical benefit (P=0.015, HR 1.481, 95%CI 1.081-2.030; P=0.003, HR 1.824, HR 1.223-2.720; P=0.001, HR 4.996, 95%CI 3.189-7.826). We also did not find the association between EGFR mutation and PFS (P=0.368). EGFR mutations mainly consist of the exon 19 deletion (del19) and the L858R point mutation. The other mutations are not as common.

We used multivariate Cox regression analysis to evaluate the independent prognostic predictors. Cox regression analysis demonstrated that NLR and, primary tumor location were related to PFS and can be considered as independent factors for predicting poor prognosis (P=0.001, HR=3.297, 95%CI= 1.614-6.737; P=0.001, HR=2.021, 95%CI=1.365-2.993; respectively) (Fig.3).

Discussion

Systemic inflammation plays a critical role in tumor proliferation and metastasis. Our study demonstrated that pre-treatment NLR (\geq 4.965) and primary tumor location were independently and significantly associated with the shorter PFS in patients with EGFR-mutated metastatic NSCLC. In univariate analyses, NLR, Lym, Lym%, Neu, and Neu% all played an important role. In addition to the NLR, previously determined related and effective prognostic factors include tumor size, sex, disease status, tumour location and performance status [2]. As the NLR as an inflammatory marker plays an increasingly role, it is widely analyzed in the solid tumors, such as ovarian cancer, urothelial carcinoma, head and neck cancer, lung cancer, hepatocellular carcinoma and so on[16]. The majority of the analyses focus on the multiple comprehensive treatments, including chemotherapy, chemoradiotherapy, radiation therapy, surgery, immunotherapy and immunotherapy combined with radiotherapy[13, 17-22]. The mechanism reflects the patients' inflammatory and systemic immune status. However, in a subset of EGFR-mutant advanced disease, NLR was an important factor to assess the prognosis when treated with chemotherapy as first-line therapy. There are few studies on patients treated with targeted therapies,

particularly EGFR-TKIs. Only a few studies have shown that the NLR is a significant prognostic factor for PFS in patients who receive TKIs. Our study complements the current studies in this field [15, 23].

To the best of our knowledge, inflammation can be regarded as the hallmark of cancer, and plays an integral role in tumorigenesis, lymphomagenesis and progression of cancer. Increasing evidence showed that elevated inflammation was related to poor cancer-specific in a variety of tumors[16, 24]. Tumor cells can lead to upregulation of the inflammatory process, which can release the proinflammatory factors, promoting the cancer cell proliferation, angiogenesis and lymphagionesis. The inflammatory cells and factors, including the lymphocytes, neutrophils, platelet, IL-6, IL-8 and C-reactive protein (CRP) have different influences in various cancers [25-27]. Neutrophils and macrophages can secrete tumor growth factors, like the TL-4, IL-8 and vascular endothelial growth factor (VEGF), which can stimulate the tumor microenvironment. Yosuke Morizawa et al analyzed the correlation of tumor microenvironment and NLR in blood samples with muscle bladder cancer, particularly in immune cells and cytokines. They found that preoperative NLR was associated with immunohistochemical expression of forkhead box P3 (Foxp3) in bladder cancer. In addition, they also suggest that IL-6 and IL-8 produced by cancer cells influence the level of NLR in patients with bladder cancer[26]. The same conclusion can also be found in head and neck squamous cell carcinoma, Ming-Shao Tsai et al demonstrated that NLR was positively related to the expression levels of IL-6 and PD-L1 expression[27]. Lymphocyte, especially the tumor-infiltrating lymphocytes (TILs) have a significant effect not only on the lung cancer but also on other solid tumors. The relationship between peripheral blood counts and CD8+ TILs has been found in breast cancer[28].

In NSCLC, TILs play a significant role in the response to anti- PD-1 therapy in patients with metastatic. The more activated CD8+ cells there are, the better the tumor can be controlled by cytotoxic activity and by inducing apoptosis of cancer cells [29]. Lymphocyte counts are also used to assess the prognosis in many lung tumors. In a recent analysis, the cut-off value for treatment induction was $\geq 1,000$ cells/ μ l to evaluate the clinical benefits treated with immunotherapy combining (RT) [30]. Preoperative lymphocyte counts are considered to be favorable prognostic factors in NSCLC to predict the disease-free survival[29]. In our study, we found that Lym% play a significant role in the PFS. The higher the elevated relative lymphocyte counts are, the better the clinical benefits are in patients receiving first-, second and third-line TKIs.

In the inflammatory response to cancer, neutrophils may play a role as reservoirs for circulating vascular endothelial growth factor and promote metastasis. Previous studies have shown that circulating neutrophils release various inflammatory factors, including tumor necrosis factor- α (TNF- α) and IL-6, to promote tumor progression [7]. In our study, we also found that the higher absolute neutrophil counts and relative neutrophil counts were associated with shorter PFS.

Leukocytes include lymphocytes, monocytes, neutrophils, eosinophils and basophils. NLR would be a simple, inexpensive and reliable pre-treatment prognostic factor for patients treated with TKIs. Iseki et al showed that Lym% was affected by neutrophils and monocytes, which is the reason why Lym% reflects systemic inflammation more accurately than Lym[31]. The results are consistent with our conclusions

from univariate analysis. Previous analyses shows that NLR can be used as an independent prognostic factor when patients receive gefitinib or erlotinib as a first-line or second-line therapy [22]. Multivariate Cox regression analyses showed that a higher pre-treatment NLR was associated with worse PFS. Besides, univariate analysis demonstrated that a LNLR at baseline was associated with a better prognosis in EGFR-mutated metastatic NSCLC. Further prospective investigations with adequate samples are needed to fully understand the prognostic value of the pre-treatment NLR.

The results from our retrospective study supports those from previous studies, showing that the NLR is a significant factor for prognosis of NSCLC. Additionally, our reports are the first to demonstrate that NLR and primary location can both be regarded as important prognostic factors in EGFR-mutated advanced NSCLC as first-, second-, and third-line therapies. An increasing number of findings have shown that primary tumor location is one of the determining factors for choosing the optimal treatment for and estimating the prognosis of patients with an advanced tumor. In our study, we used definitions according to previous findings in CT scans and bronchoscopies. A high portion of patients with peripheral adenocarcinoma show clinical benefit compared to patients with central adenocarcinoma. Wang et al has investigated that central adenocarcinoma has a worse prognosis compared with peripheral adenocarcinoma, which consistent with our conclusions[6]. EGFR mutation status can also be considered as a prognostic factor for treatment of TKIs as a first-line treatment in advanced NSCLC. According to a previous analysis, patients with an EGFR del19 had longer PFS than those with EGFR exon 21 mutation. Jiang et al concluded that EGFR mutation status is a good predictor for patients treated with EGFR-TKIs in NSCLC[32]. A large meta-analysis on patients who were treated with first-line TKIs also revealed that patients with an EGFR exon 21 mutation had a shorter PFS than patients with an EGFR del19[33]. However, in our research, we compared three different EGFR statuses and did not have the same conclusions. We concluded that the NLR and primary tumor locations are both predictive factors for the efficacy of EGFR-TKIs as first-, second- and third-line therapies.

We are aware that there are some limitations in our analysis. First, as a retrospective study, we have some selection biases. Although patients' data concerning the laboratory, CT scans/PET-CT and survival data were complete, there was also a patients' selection bias. Second, the relative numbers of eligible patients were small. In summary, the Lym, LLym%, HNeu, HNeu% and higher HNLR, correlated with poorer prognosis of NSCLC patients treated with TKIs. The NLR and peripheral-type tumor seem clinically meaningful for patients treated with EGFR-TKIs. As an effective and prognostic biomarker, NLR is inexpensive and readily available. We need further investigations with a large prospective study to validate our results in the future.

Conclusion

The NLR and primary tumor location are both effective and meaningful factors for EGFR-mutated advanced lung cancer (Fig3). The primary tumor location is also a significant predictor for treatment planning. We also found that NLR was a useful predictor for systemic inflammation in patients treated with TKIs. However, we need more data to explore understand the relationship among these parameters.

Our findings support the existing hypothesis that systemic inflammation is associated with clinical outcomes.

Abbreviations

EGFR: epidermal growth factor receptor; TKIs: tyrosine kinase inhibitors; NLR: neutrophil-to-lymphocyte ratio; PFS: progression free survival; NSCLC: non-small cell lung cancer; Lym: absolute lymphocyte count; Lym%: percentage of lymphocytes; Neu: absolute neutrophil count; Neu%: the percentage of neutrophil granulocytes; LMR: lymphocyte-to-monocyte ratio; ICIs: immune checkpoint inhibitors; OS: overall survival; CBC: complete blood count; CT: Computed tomography; PET: position emission tomography; PNA: peptide nucleic acid; PCR: Polymerase Chain Reaction; NGS: next generation sequencing; ROC: receiver operating characteristic; AUC: Area Under the Curve; HLym: higher in absolute lymphocyte counts; HLym%: higher in relative lymphocyte counts; LLym: lower in absolute lymphocyte counts; LLym%: lower in relative lymphocyte counts; HNeu: higher in absolute neutrophil count; HNeu%: higher in the percentage of neutrophil granulocytes; HNLR: higher in NLR; LNEU: lower in absolute neutrophil count; LNeu%: lower in the percentage of neutrophil granulocytes; LNLR: lower in NLR.

Declarations

Ethics approval and consent to participate. The studies involving human participants were reviewed and approved by The Ethics Committee of Shandong Cancer Hospital Affiliated to Shandong First Medical University. The patients/participants provided agree to participate in this study.

Consent for publication

Not applicable.

Availability of data and materials

All the data and material supporting the findings are present in the manuscript.

Competing interests

There is no conflict of interest among authors to disclose.

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Authors' contributions

Writing - original draft: XQX; Writing - review and editing: JMY, MHL and PX; Conceptualization: PX, XLL, WJT; Data collection and analysis: WJT, JLC; All authors have read and approved the final manuscript.

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Tables

| Table 1 Patients and clinical characteristics | |
|---|-------------------------------|
| Characteristics | All patients (n=190) n (%) |
| Gender | |
| Male | 68(35.8%) |
| Female | 122(64.2%) |
| Age | |
| Median | 58 |
| Range | 34-87 |
| Smoking habits | |
| Non-smoker | 148(77.9%) |
| Ever-smoker | 42(22.1%) |
| EGFR-mutation | |
| L858R | 110(57.9%) |
| 19del | 75(39.5%) |
| Rare mutation | 5(2.6%) |
| Stage | |
| Postoperative recurrence | 26(13.7%) |
| IV | 164(86.3%) |
| EGFR Mutation Test | |
| PCR | 172(90.5%) |
| NGS | 18(9.5%) |
| Tumor location | |
| Central type | 48(25.3%) |
| Peripheral type | 142(74.7%) |
| Bone metastasis | |
| YES | 83(43.7%) |
| NO | 107(56.3%) |
| Liver metastasis | |
| YES | 26(13.7%) |
| NO | 164(86.3%) |
| Brain metastasis | |
| YES | 59(31.1%) |
| NO | 131(68.9%) |
| First-line therapy | |
| Surgery | 27(14.2%) |
| Chemotherapy | 93(49.0%) |
| Chemoradiotherapy | 9(4.7%) |
| Radiotherapy combined targeted therapy | 15(7.9%) |
| Targeted therapy | 36(18.9%) |
| Radiotherapy | 6(3.2%) |
| Chemotherapy combined with TKIs | 4(2.1%) |
| Target therapy lines | |
| First-line | 54(28.4%) |
| Second-line | 102(53.7%) |
| Third-line | 34(17.9%) |

Table 2

Univariate and multivariate Cox proportional analysis regarding PFS (n=190)

| Variate | Univariate analyses | | Multivariate analyses | |
|--------------------------|---------------------|---------------------|-----------------------|---------------------|
| | P | HR (95% CI) | P | HR (95% CI) |
| Age(years) | 0.387 | 1.007(0.991- 1.024) | 0.735 | 1.003(0.986-1.021) |
| Gender | | | | |
| Male | | | | |
| Female | 0.725 | 0.943(0.681-1.306) | 0.521 | 0.874(0.579-1.318) |
| Smoking Habit | | | | |
| Non-smoker | | | | |
| Ever-smoker | 0.620 | 1.100(0.755-1.602) | 0.794 | 0.935(0.563-1.551) |
| EGFR Mutation | | | | |
| L858R | | | | |
| 19del | | | | |
| Rare mutation | 0.239 | 1.198(0.887-1.618) | 0.192 | 1.262(0.890-1.789) |
| EGFR Mutation test | | | | |
| PCR | | | | |
| NGS | 0.291 | 0.726(0.400-1.315) | 0.613 | 1.190 (0.606-2.340) |
| Stage | | | | |
| Postoperative recurrence | | | | |
| IV | 0.934 | 0.981(0.630-1.5280) | 0.655 | 1.414(0.639-2.040) |
| Location | | | | |
| Central | | | | |
| Peripheral | 0.003 | 1.739(1.207-2.506) | 0.000 | 2.021(1.365-2.993) |
| Metastatic lesions | | | | |
| Bone | 0.324 | 1.172 (0.855-1.607) | 0.924 | 0.982(0.682-1.415) |
| Liver | 0.160 | 1.417(0.872-2.302) | 0.856 | 1.052(0.608-1.820) |
| CNS | 0.955 | 1.010(0.725-1.406) | 0.952 | 0.988(0.673-1.451) |
| Lym ($\times 10^9$) | | | | |
| <1.625 | | | | |
| ≥ 1.625 | 0.012 | 0.662(0.480-0.915) | 0.895 | 0.971(0.628-1.502) |
| Lym% | | | | |
| <18.8% | | | | |
| $\geq 18.8\%$ | 0.000 | 0.391(0.274-0.560) | 0.117 | 0.625(0.347-1.125) |
| Neu ($\times 10^9$) | | | | |
| <3.675 | | | | |
| ≥ 3.675 | 0.015 | 1.481(1.081-2.030) | 0.766 | 1.068(0.691-1.651) |
| Neu% | | | | |
| <51.8% | | | | |
| $\geq 51.8\%$ | 0.003 | 1.824(1.223-2.720) | 0.079 | 1.582(0.949-2.638) |
| NLR | | | | |
| <4.965 | | | | |
| ≥ 4.965 | 0.000 | 4.996(3.189-7.826) | 0.001 | 3.297(1.614-6.734) |
| First-line therapy | 0.409 | 1.048(0.983-1.127) | 0.665 | 0.969(0.841-1.117) |
| Target therapy lines | 0.782 | 0.968(0.767-1.221) | 0.460 | 1.114(0.837-1.483) |

Table 3 The optimal cut-off values based on PFS

| Peripheral blood index | Sensitivity | Specificity | AUC |
|-------------------------|-------------|-------------|-------|
| Lym ($\times 10^9$ /L) | 63% | 64.4% | 0.632 |
| Lym% | 88.9% | 28.2% | 0.549 |
| Neu ($\times 10^9$ /L) | 59.3% | 54% | 0.521 |
| Neu% | 88.9% | 19% | 0.499 |
| NLR | 17.8% | 100.0% | 0.550 |

Table 4 Peripheral blood test with clinicopathological factors with P values

| Parameter | R spearman | P value | U test | P value |
|-----------------------|------------|---------|----------|---------|
| NLR& Lym | -0.310 | 0.000 | | |
| NLR& Neu | 0.595 | 0.000 | | |
| Lym%& Bone metastatic | | | 3617.500 | 0.029 |
| Neu%& Bone metastatic | | | 5442.5 | 0.008 |
| NLR& Bone metastatic | | | 5299.5 | 0.022 |

Figures

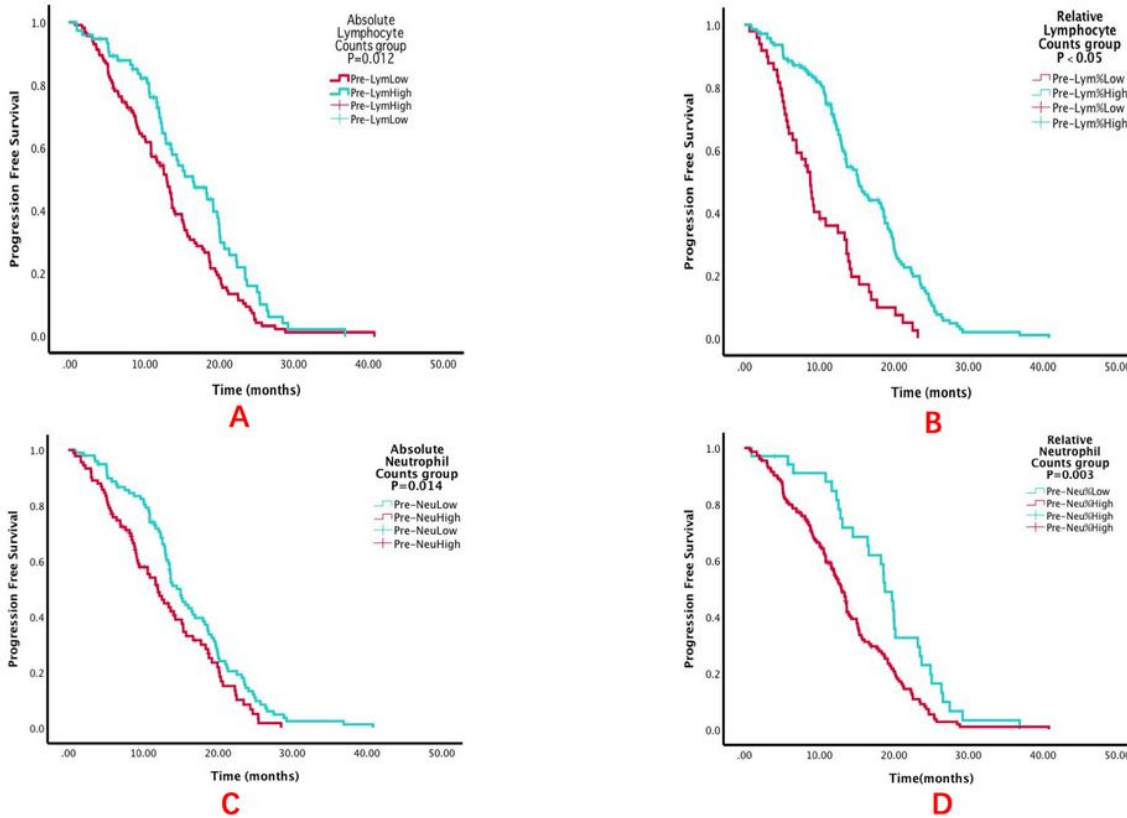


Figure 1

Kaplan-Meier plot of progression-free survival (PFS) stratified by pretreatment absolute lymphocyte counts group (A), relative lymphocyte counts group (B), absolute neutrophil counts group (C), and relative neutrophil counts group (D).

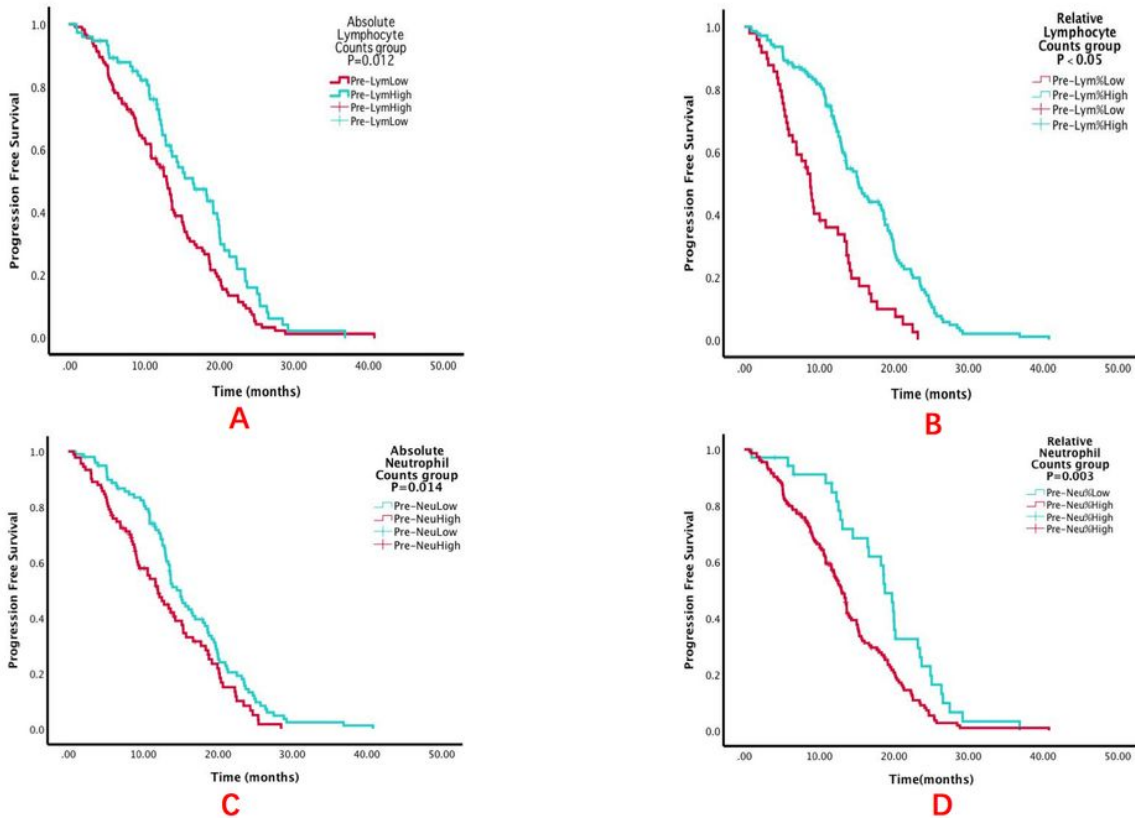


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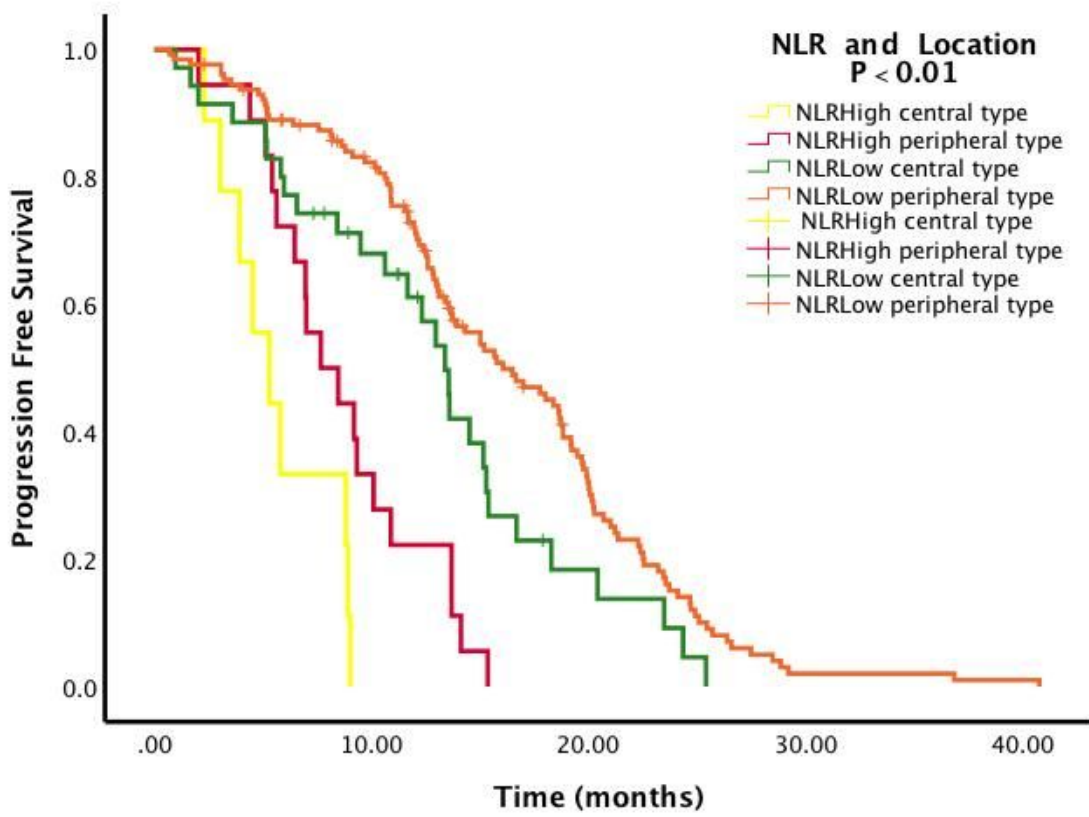


Figure 2

Kaplan-Meier survival curves of PFS based on combination of pretreatment neutrophil to lymphocyte expression and primary location in all patients.

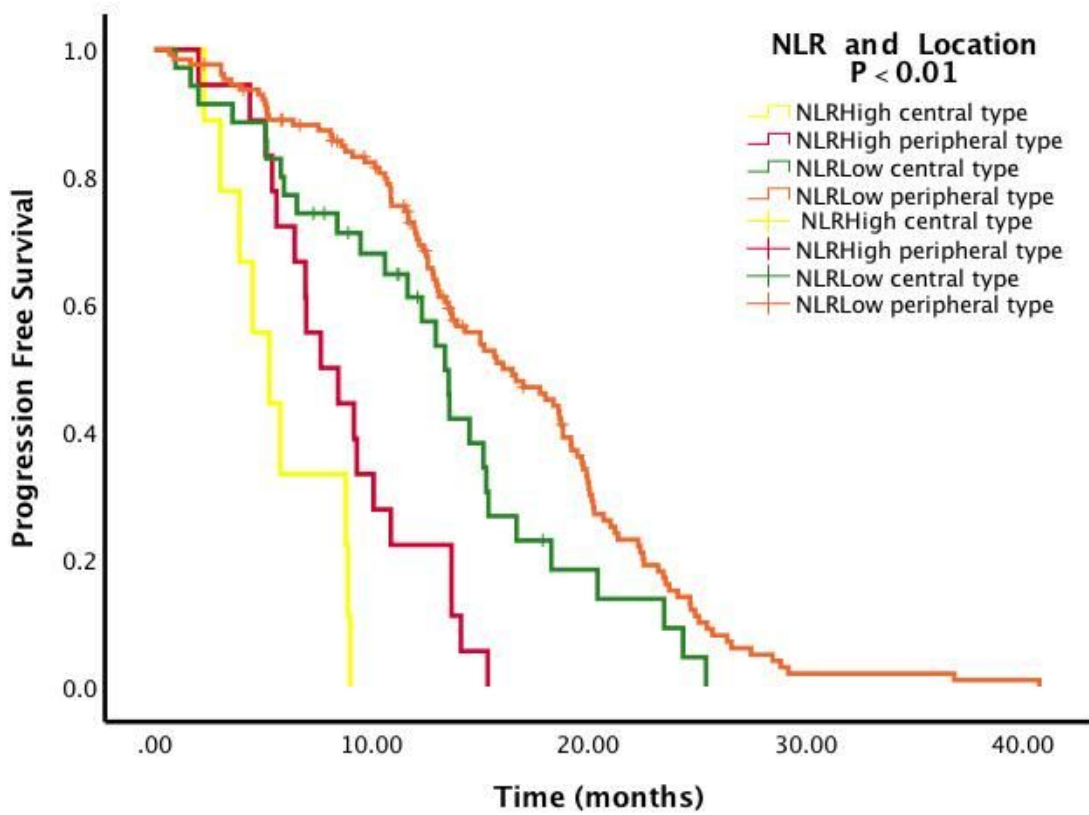


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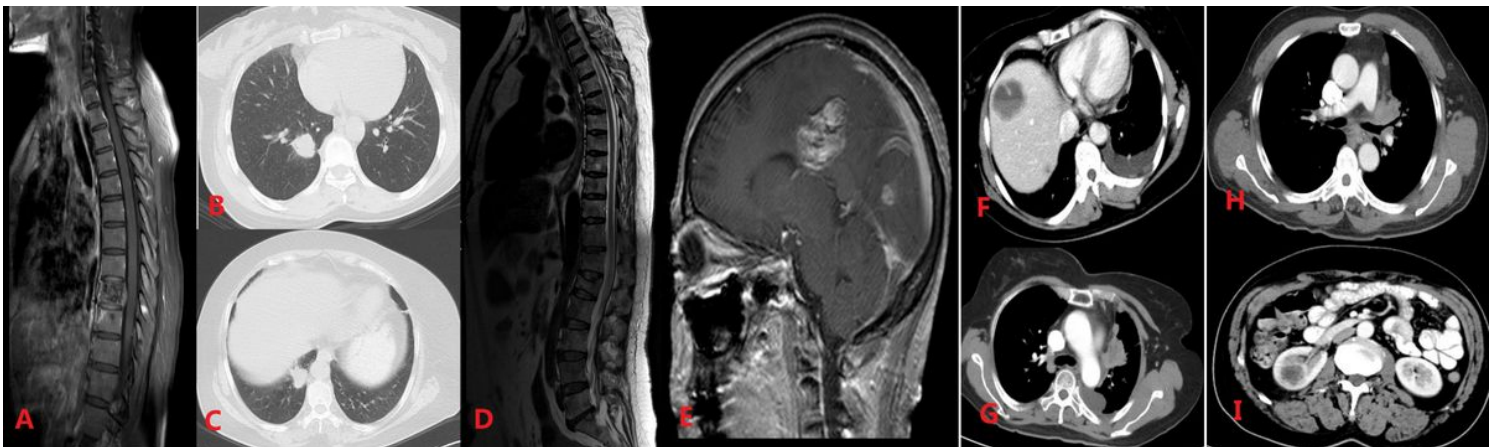


Figure 3

A patient with high NLR and peripheral type (B) in stage IV non-small cell lung cancer (A) with shorter PFS. A patient with low NLR and peripheral type (C) in stage IV lung adenocarcinoma with longer PFS. (D) A patient with low NLR and central (G) in stage IV lung adenocarcinoma (E, F) with shorter PFS. A patient with high NLR and central type (H) in stage IV (I) with longer PFS.

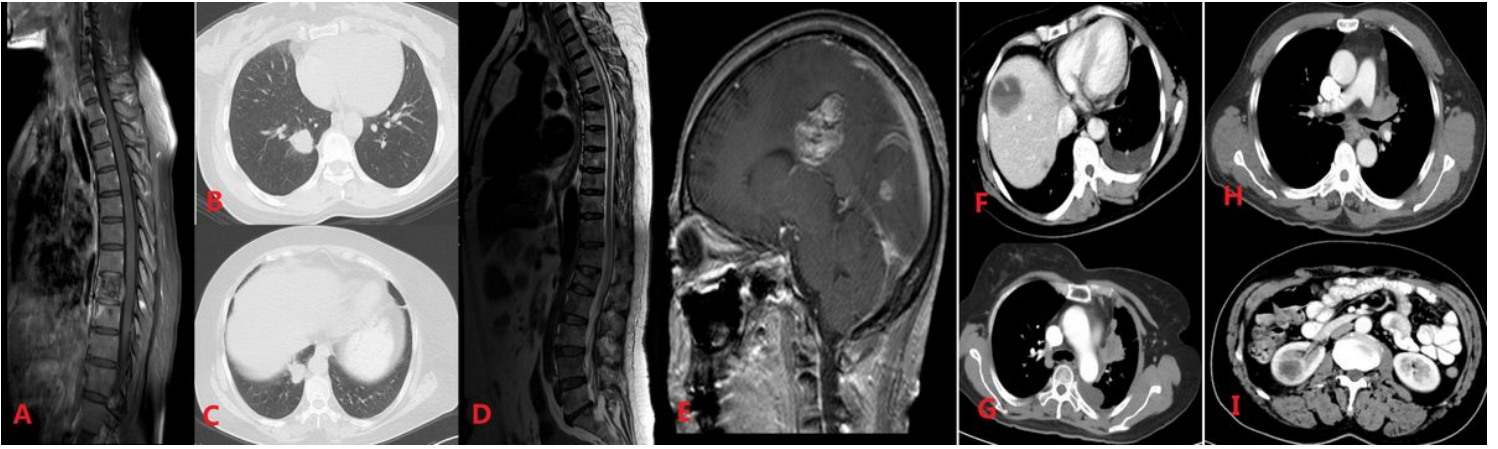


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