

Association between CD4⁺ T cells ATP levels and disease progression in patients with non-small cell lung cancer

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Abstract. Introducing the exploration of stimulated CD4⁺ cells adenosine triphosphate (sATP^{CD4}) levels for immune monitoring post non-small cell lung cancer (NSCLC) chemotherapy, the present study aimed to investigate its efficacy in gauging the potential risk of disease progression (PD) in patients with NSCLC. Therefore, a total of 89 patients with advanced NSCLC, who underwent chemotherapy between August 15 2022 and August 30 2023 at the Fifth Affiliated Hospital of Guangzhou Medical University (Guangzhou, China), were retrospectively studied. Patients were divided into the PD (n=21) and disease stability (non-PD; n=68) groups and their clinical data were compared. The thresholds for predicting PD were identified using receiver operating characteristics (ROC) curves. Multivariate logistic regression analysis was carried out to assess the association between peripheral blood markers and the incidence of PD. Therefore, post-chemotherapy, significant differences in white blood cell count, non-stimulated CD4⁺ cells ATP and sATP^{CD4} levels were obtained between patients in the PD and non-PD groups (P<0.05). In addition, sATP^{CD4} levels were notably decreased in the PD group compared with the non-PD group. Furthermore, ROC analysis revealed that the predictive threshold for PD was 224.5 ng/ml [area under the curve=0.887; 95% confidence interval, 0.811-0.963]. Additionally, patients with low immunity (ATP <224.5 ng/ml) exhibited a higher risk of PD compared with the high-immunity group (ATP >224.5 ng/ml; P<0.0001). Finally, multivariate logistic regression analysis suggested that sATP^{CD4} could serve as an independent factor for predicting NSCLC progression. Overall, the current study

predicted that immune function could be possibly associated with the risk of PD in patients with NSCLC.

Introduction

Lung cancer, a major factor highly associated with cancer-related mortality worldwide, resulted in 710,000 deaths in China alone, out of 3 million cancer-related fatalities in 2020. Non-small cell lung cancer (NSCLC) is involved in ~80% of these cases (1,2).

Notably, the majority of these patients are diagnosed at an advanced stage of the disease, making them unsuitable candidates for surgical resection. Advanced-stage lung cancer patients are commonly treated with platinum-based chemotherapy regimens, which exert favorable efficacy (3). However, in addition to the therapeutic benefits of platinum-based regimens, attention should be also paid to their hematologic and gastrointestinal toxicities, which can potentially significantly compromise the patient's immune function and treatment outcomes (4). According to the cancer immunoeediting theory, including immune elimination, homeostasis and escape, indicates that immunity is closely associated with the occurrence and progression of cancer (5). Therefore, assessing and improving the immune function of late-stage patients with NSCLC following chemotherapy has become a clinical concern.

Previous studies indicated that compared with healthy individuals, patients with lung cancer exhibited lower count of CD4⁺ T and NK cells and reduced CD4⁺/CD8⁺ ratios, accompanied by elevated number of regulatory T cells (Tregs) (6,7). Additionally, another study showed that patients with NSCLC, who responded effectively to immunotherapy, experienced an increase in the proportion of CD4⁺ T cells compared with the baseline values, while no statistically significant changes were observed in the ineffective response group (8). Furthermore, previous studies also demonstrated that chemotherapy drugs could impair the proliferation and function of peripheral circulating effector T cells, thus potentially leading to the reduced proliferative activity of cytotoxic T lymphocytes (CTLs) (9,10). However, it has been also reported that chemotherapy drugs can induce tumor cell apoptosis, which in turn trigger immune responses, eventually enhancing the cytotoxicity of CTLs and strengthening anti-tumor immunity (11). Chemotherapy drugs can also accelerate CD4⁺CD25⁺ Treg apoptosis, thus

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reducing the number of Tregs and effectively regulating tumor immunity (12). In summary, there is no consensus regarding the effect of chemotherapy on the immune function of patients with cancer and the immune status on chemotherapy outcome. Therefore, more research and immune function assessment methods are needed to establish a conclusive understanding.

The CD4⁺ T cell adenosine triphosphate (ATP) release assay is used to measure intracellular ATP concentration within purified CD4⁺ T lymphocytes following stimulation with phytohemagglutinin-L. This assay is used to assess peripheral immune function and emerging evidence suggests that it has significant predictive value in predicting post-transplant infections in organ transplant recipients and hematopoietic stem cell recipients, as well as septic shock in septic patients (13-17).

Some researchers have found that patients who developed septic shock had markedly lower CD4⁺ T cell ATP levels compared with septic patients who did not shock. However, there were no significant differences in the percentages of CD4⁺ and CD8⁺ T lymphocytes between the two groups (17). Additionally, Serban *et al* (16) conducted a study showed that patients with decreased CD4⁺ T cell ATP levels five months after allogeneic transplantation had a higher risk of infection, while CD4⁺ T cell counts remained at an increased level (16). This indicates that for patients with similar immune cell quantities and poorer health status sATP^{CD4} holds a special clinical value. Clinical trials have been conducted to evaluate CD4⁺ T cells ATP release levels in hepatocellular carcinoma, thus revealing a close association between progression-free survival (PFS), overall survival (OS) and cancer recurrence (18,19). Tumor initiation and progression are intricately associated with the host immune function. Tumor immunity represents a dynamic process, where T cell responses are enhanced. Therefore, maintaining adequate immune function is crucial for fostering a beneficial cycle (20). However, to the best of our knowledge, the association between CD4⁺ T cells ATP levels and NSCLC progression has not been previously investigated. Therefore, exploring the association between CD4⁺ T cells ATP release levels and NSCLC holds potential value in understanding the interplay between immune function and NSCLC progression.

The present study aimed to evaluate the changes in immune function prior and after chemotherapy, as well as the association between post-chemotherapy immune status and tumor progression.

Materials and methods

Patients. Patients diagnosed with NSCLC, who were treated with chemotherapy, were retrospectively reviewed between August 15 2022 and August 30 2023, at the Fifth Affiliated Hospital of Guangzhou Medical University Hospital, Guangzhou, China (n=183). Among a total of 183 patients, 89 patients were screened and allocated into the disease progression (PD, n=21) and disease stability (non-PD; n=68; Fig. 1) groups, males accounted for 52.8% of the population, with patient ages ranging from 33 to 78 years (mean age 66.2±10.2). A total of 10 and 20 patients in the PD and non-PD groups, respectively, underwent immunocellular function assay before chemotherapy. The present study received approval from the Ethics Committee of the Fifth Affiliated Hospital of Guangzhou Medical University (Guangzhou, China) dated

08-07-2022 of and approval no. KY01-2022-07-08. Written informed consent was obtained from all participants, allowing the use of their clinical data in this study. The research adhered to the principles outlined in the Declaration of Helsinki.

Inclusion and exclusion criteria. The inclusion criteria were as follows: i) Patients with pathological diagnosis of NSCLC; ii) with locally advanced disease, medically or technically inoperable; iii) those treated with periodic chemotherapy (pemetrexed, cisplatin, carboplatin, paclitaxel, gemcitabine); iv) with Eastern Cooperative Oncology Group (ECOG) score of 0-2 (ECOG score 0: Normal activity without symptoms, fully active and able to carry on all pre-disease performance without restriction; ECOG score 1: Symptomatic, but completely ambulatory, restricted in physically strenuous activity, but able to perform light or sedentary work; ECOG score 2: Symptomatic, in bed <50% of waking hours and capable of limited self-care and confined to bed or chair >50% of waking hours) (21). Patients were staged according to the 8th edition TNM staging system (22). The exclusion criteria were the followings: i) Patients with stage I, II or III NSCLC; ii) those who were treated with immunotherapy; iii) HIV-positive individuals; and iv) those whose ATP release levels from CD4⁺ T lymphocytes were not recorded after four cycles of chemotherapy. The study conformed to the principles outlined in the Declaration of Helsinki (approval no. KY01-2022-07-08).

Study assessment. Neck, chest, whole abdomen and pelvic computed tomography (CT) imaging examinations were conducted prior to initial treatment and at 20 days following the completion of four cycles of treatment. Treatment efficacy was evaluated according to the Response Evaluation Criteria for Solid Tumors (RECIST) 1.1 criteria (23) and categorized as PR (partial response), SD (stable disease) or PD. Patients who were treated with chemotherapy for four cycles and had no progression at two consecutive radiological assessments were allocated into the non-PD group (PR + SD), while those who did not meet the aforementioned criteria were included in the PD group. The current analysis was performed to assess the association between peripheral blood CD4⁺T cells ATP expression and treatment efficacy after four cycles of therapy.

Immunocellular function assay. The levels of ATP released by stimulated (sATP^{CD4}) and non-stimulated (nATP^{CD4}) CD4 T lymphocytes were assessed through a luciferin-luciferase reaction, according to the manufacturer's instructions (AIMdex[®]; Leide Biosciences Co., Ltd.) (24). Briefly, 100 μ l of 4-fold diluted whole blood was incubated with 8.75 ng/ml sATP or without phytohaemagglutinin (PHA; nATP) for 15-18 h, 37°C. CD4⁺ T cells were isolated using magnetic beads (900-1, 100 μ g/well; Leide Biosciences Co., Ltd.) and ATP was released using a lysis buffer. Subsequently, the mixture was supplemented with luciferin/luciferase and bioluminescence was measured using a luminometer (JR-I; Weihai Weigao Biotechnology Co., Ltd). The results were analyzed using the corresponding software (JR-1 v2.3.8.8.181120/1.2.0; Weihai Weigao Biotechnology Co., Ltd). A standard curve with ATP calibrators (0, 50, 100, 200, 400 and 800 ng/ml) was constructed. ATP release was determined by comparing ATP levels between CD4⁺T cell- stimulated and non-stimulated

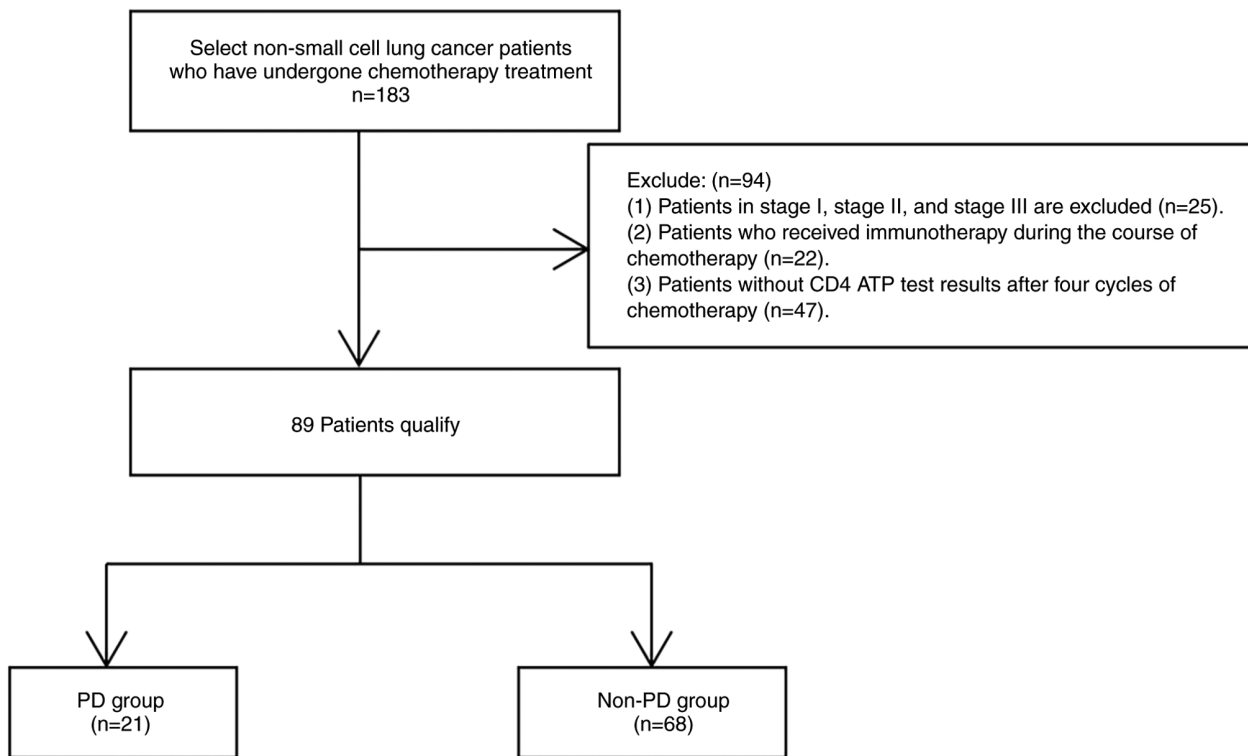


Figure 1. Flow chart of the study population. A retrospective analysis was conducted on patients with non-small cell lung cancer, who underwent chemotherapy at the Fifth Affiliated Hospital of Guangzhou Medical University (Guangzhou, China) between August 15, 2022 and August 30, 2023 (n=183). Based on strict inclusion and exclusion criteria, a total of 89 patients were ultimately included and divided into the disease progression (n=21) and disease stability (n=68) groups. ATP, adenosine triphosphate; PD, disease progression.

samples (Fig. 2) and validated through clinical repeatability assessment, meeting clinical standards.

Blood parameter assessment. White blood cell (WBC), granulocyte and lymphocyte counts were measured using a fully automated hematology analyzer (Original Mindray BC-5180 CRP; Mindray).

Treatment and data collection. The relevant clinical and laboratory data were collected from the electronic medical records system. The clinical characteristics of patients, such as age, sex, smoking and drinking history, underlying medical conditions and histology were collected and summarized (Table I). Baseline measurements were defined as those after chemotherapy.

Statistical analysis. All statistical analyses were performed using SPSS 26.0 software (IBM Corp.). The clinical and demographic data are expressed as the mean \pm standard deviation (SD) or medians [interquartile ranges (IQRs)] for continuous variables, depending on their distribution (normally and non-normally distributed). Categorical variables are expressed as percentages. The differences between categorical, normally distributed and non-normally distributed continuous variables were compared using chi-square, Student's t-test (Between-group comparisons were performed using independent t-tests, while within-group comparisons before and after treatment were conducted using paired t-tests) and Mann-Whitney U test, respectively. The association between peripheral blood biomarkers and PD was assessed

by logistic regression analysis, while the predictive values of the baseline peripheral blood parameters were evaluated using receiver operating characteristics (ROC) curves. In addition, the optimal cut-off values were determined using the Youden's index, which was calculated using the following formula: Youden's index=Sensitivity + Specificity-1. The maximum Youden's index value was considered to indicate the optimal cut-off point. The differences in area under the curve (AUC) values were analyzed using DeLong test. All graphs were generated using GraphPad Prism 8.0 software (Dotmatics).

Results

Patient characteristics. In the present study, a total of 89 patients were allocated into the non-PD (n=68; including three PR and 65 SD cases) and PD (n=21 cases) groups. The baseline characteristics of patients are summarized in Table I. No significant differences were recorded in terms of sex (P=0.296; $\chi^2=1.092$), age (P=0.174; Z=0.119), body mass index (BMI; P=0.258; t=1.178), smoking history (P=0.867; $\chi^2=0.028$), drinking history (P=0.747; $\chi^2=0.104$), diabetes (P=0.747; $\chi^2=0.104$), hypertensive (P=0.304; $\chi^2=1.057$), histology (P=0.561; $\chi^2=0.228$), distant metastasis (P=0.962; $\chi^2=0.002$), EGFR mutation status (P=0.052; $\chi^2=3.760$), treatment regimen (P=0.829; $\chi^2=0.884$) and chemotherapy regimen (P=0.125; $\chi^2=4.160$) between the PD and non-PD groups.

Comparison of peripheral blood indicators between the PD and non-PD groups. Peripheral blood indicators were compared between the PD and non-PD groups (Fig. 3). The

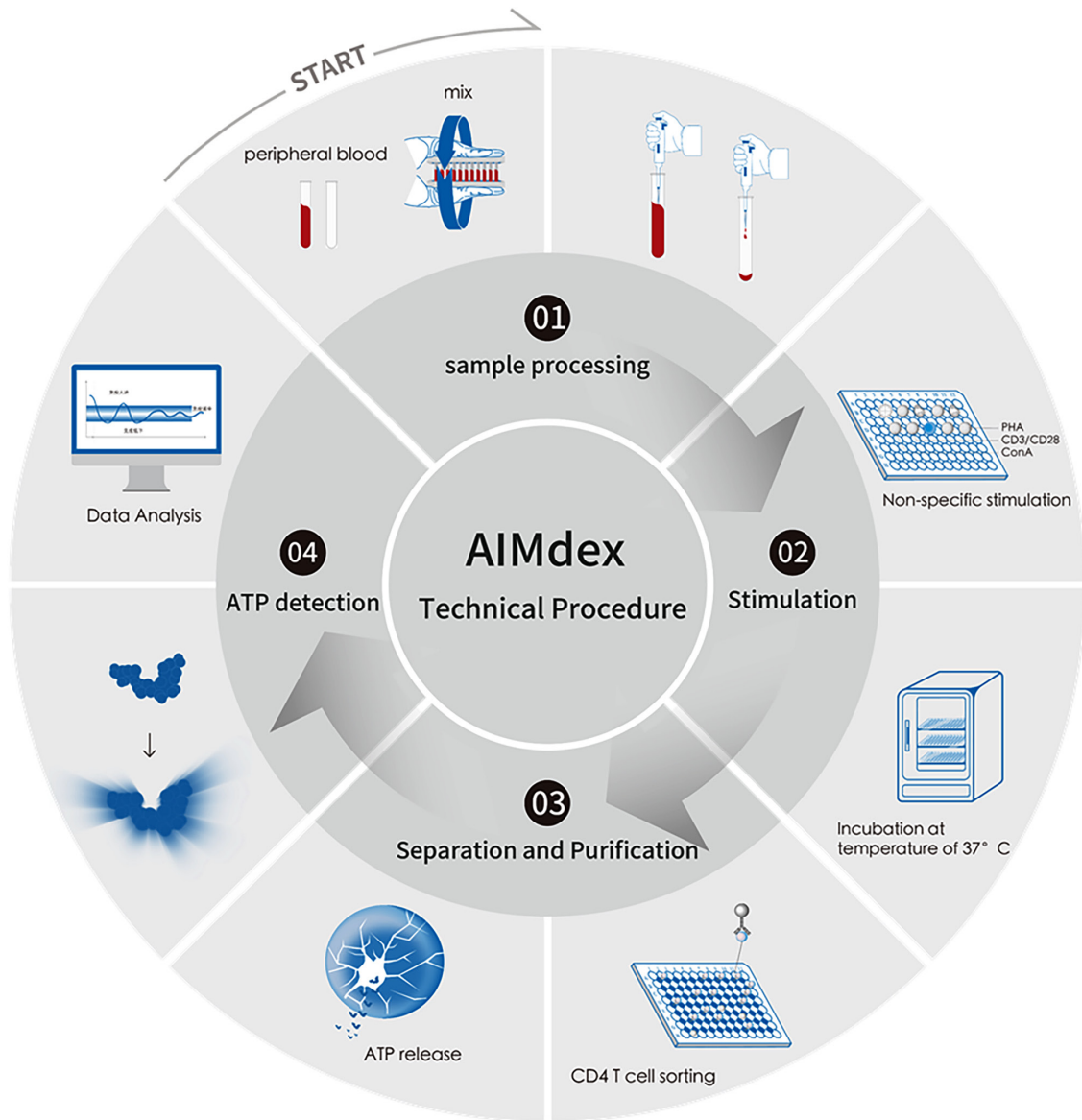


Figure 2. Immunocellular function assessment workflow (Leide Biosciences Co., Ltd. was the source of the flowchart, used with permission). The Immunocellular function assessment process was divided into the following four steps: Step 1: Sample processing: 3-5 ml peripheral blood was collected and mixed well. Subsequently, 250 μ l peripheral blood was diluted to 1 ml and mixed well; Step 2: Stimulation: Each blood sample was seeded into a 96-well cell culture plate (one well for non-stimulated and one well for phytohaemagglutinin-stimulated cells). Cells were incubated at 37°C in an incubator for 15-18 h; Step 3: Separation and purification: Each well of the culture plate was supplemented with CD4 magnetic beads, followed by incubation at room temperature for 20 min. Subsequently, CD4⁺ T lymphocytes were isolated using a magnetic rack and lysed with a lysis buffer. Following the addition of luciferin/luciferase mixture, bioluminescence was measured utilizing a luminometer for 5 min; Step 4: ATP detection: ATP levels were analyzed using a luminometer (JR-I; Weihai Weigao Biotechnology Co., Ltd). ATP, adenosine triphosphate.

analysis revealed a significant reduction in WBC count ($P=0.028$; Fig. 3A) and $nATP^{CD4}$ ($P=0.002$; Fig. 3D) and $sATP^{CD4}$ ($P<0.001$; Fig. 3E) levels between the PD and non-PD groups (Fig. 3A). Additionally, a slight decrease in neutrophil ($P=0.059$; Fig. 3B) and lymphocyte counts ($P=0.122$; Fig. 3C) was also recorded.

Changes in the peripheral blood indicators after chemotherapy. To comprehensively assess the alterations in peripheral blood indicators following chemotherapy, the levels of the relevant indicators prior and after four cycles of treatment were compared. Data from both the pre- and post-treatment periods were available for 10 PD and

20 non-PD cases. No significant changes in WBC, neutrophil and lymphocyte counts and $nATP^{CD4}$ levels were obtained after chemotherapy in both the PD and non-PD groups ($P>0.05$; Fig. 4A-H). However, $sATP^{CD4}$ levels were notably reduced in the PD group after chemotherapy ($P=0.008$; Fig 4I), while they were significantly enhanced in the non-PD group ($P<0.0001$; Fig 4J).

Cut-off points and association with the occurrence of PD. The diagnostic efficacy of WBC, $nATP^{CD4}$ and $sATP^{CD4}$ after four treatment cycles was assessed by ROC curves (Fig. 5). For $sATP^{CD4}$, a cut-off value of 224.5 ng/ml [AUC=0.887; 95% confidence interval (CI), 0.811-0.963; specificity 88.2%,

Table I. Comparison of baseline characteristics between PD group and non-PD group.

Variables	PD (n=21)	non-PD (n=68)	$\chi^2/Z/t$	P-value
Sex			1.092	0.296
Male, n (%)	9 (42.9)	38 (55.9)		
Female, n (%)	12 (57.1)	30 (44.1)		
Age, years (interquartile Range)	64.0 (54.0-71.5)	67.5 (57.0-72.0)	0.119	0.174
BMI, (kg/m ²)	22.4±3.7	21.4±3.1	1.178	0.258
Smoking history			0.028	0.867
Yes, n (%)	5 (23.8)	15 (22.1)		
pack-year (≥ 30), n (%)	5 (100)	15 (100)		
No, n (%)	16 (76.2)	53 (77.9)		
Drinking history			0.104	0.747
Yes, n (%)	2 (9.5)	5 (7.4)		
Every day, n (%)	2 (100)	5 (100)		
No, n (%)	19 (90.5)	63 (92.6)		
Diabetes			0.104	0.747
Yes, n (%)	2 (9.5)	5 (7.4)		
Type1	0	0		
Type2, n (%)	2 (100)	5 (100)		
No, n (%)	19 (90.5)	63 (92.6)		
Hypertensive			1.057	0.304
Yes, n (%)	3 (14.3)	17 (25.0)		
Stage 1, n (%)	1 (33.3)	2 (11.8)		
Stage 2, n (%)	2 (66.7)	10 (58.8)		
Stage 3, n (%)	0	5 (29.4)		
No, n (%)	18 (85.7)	51 (75.0)		
Histology			0.338	0.561
Non squamous carcinoma, n (%)	19 (90.5)	64 (94.1)		
Squamous carcinoma, n (%)	2 (9.5)	4 (5.9)		
Distant metastasis			0.002	0.962
Yes, n (%)	18 (85.7)	58 (85.3)		
No, n (%)	3 (14.3)	10 (14.70)		
EGFR mutation			3.760	0.052
EGFR ⁺	18 (85.7)	43 (63.2)		
L858R, n (%)	3 (16.6)	6 (14.0)		
Deletion 19, n (%)	0	1 (2.3)		
EGFR18, n (%)	1 (5.6)	2 (4.6)		
EGFR20, n (%)	3 (16.6)	6 (14.0)		
EGFR21, n (%)	1 (5.6)	6 (14.0)		
Other, n (%)	10 (55.6)	22 (51.1)		
EGFR ⁻ , n (%)	3 (14.3)	25 (36.7)		
KRAS mutation			-	-
KRAS ⁺ , n (%)	2 (9.5)	5 (7.4)		
pG12C ⁺ , n (%)	2 (100)	2 (40)		
pG12D ⁺ , n (%)	0	3 (60)		
Not detected, n (%)	19 (90.5)	63 (92.6%)		
Treatment regimen			0.884	0.829
Chemotherapy only, n (%)	4 (19.0)	9 (13.2)		
Chemotherapy + anti-EGFR, n (%)	6 (28.6)	21 (30.8)		
Chemotherapy + anti-VEGF, n (%)	9 (42.9)	34 (50.0)		
Chemotherapy + anti-EGFR + anti-VEGF, n (%)	2 (9.5)	4 (5.9)		

Table I. Continued.

Variables	PD (n=21)	non-PD (n=68)	$\chi^2/Z/t$	P-value
Chemotherapy regimen			4.160	0.125
PemC/PemP, n (%)	11 (52.4)	49 (72.1)		
GP, n (%)	9 (42.9)	14 (20.6)		
TC, n (%)	1 (4.8)	5 (7.4)		
Chemotherapy + anti-EGFR + anti-VEGF, n (%)	2 (9.5)	4 (5.9)		

BMI, Body Mass Index; EGFR, epidermal growth factor receptor; VEGF, Vascular endothelial growth factor. PemC, pemetrexed + Carboplatin; PemP, pemetrexed + Cisplatin; GP, Gemcitabine + Carboplatin/Cisplatin; TC, Paclitaxel + Carboplatin/Cisplatin;

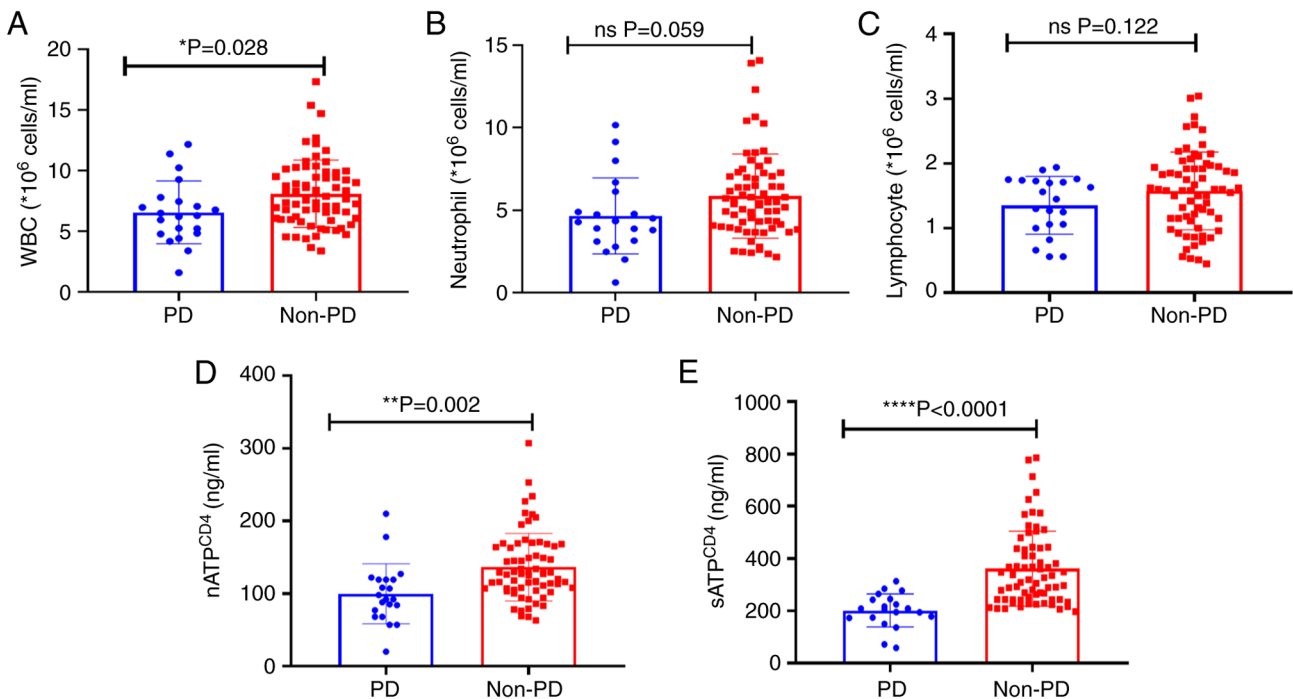


Figure 3. Comparison of peripheral blood indicators between the PD (n=21) and non-PD groups (n=68). (A) White blood cell count was significantly lower in the PD group compared with the non-PD group (P=0.028). (B) No significant difference in neutrophil count was observed between the PD and non-PD groups. (C) No significant difference in lymphocyte count was observed between both groups. (D) Non-stimulated CD4⁺ cell ATP levels in the PD group were notably decreased compared with the non-PD group (P=0.002). (E) sATP^{CD4} concentration PD and Non-PD groups. Stimulated CD4⁺ cell ATP levels in the PD group were markedly lower compared with the non-PD group (P<0.0001). *P<0.05, **P<0.01 and ****P<0.0001; ns, no significant. PD, disease progression; ATP, adenosine triphosphate.

sensitivity 71.4%] displayed the highest sensitivity and specificity for PD diagnosis. Consistently, cut-off values of 99 ng/ml (AUC=0.713; 95% CI, 0.580-0.846; specificity 82.4%, sensitivity 52.4%) and 7.09×10^9 cells/l (AUC=0.660; 95% CI, 0.524-0.796; specificity 60.3%, sensitivity 71.4%) were obtained for nATP^{CD4} and WBC, respectively. The AUC value for sATP^{CD4} was significantly higher compared with that for nATP^{CD4} (P=0.010) and WBC (P=0.001; Table II).

Analysis of risk factors associated with the occurrence of PD. Univariate and multivariate logistic regression analyses were performed to explore the association between measured variables and PD occurrence. In univariate analysis, WBC count (OR=3.796; 95% CI, 1.310-11.003; P=0.014) and

nATP^{CD4} (OR=5.133; 95% CI, 1.780-14.806; P=0.002) and sATP^{CD4} (OR=18.750; 95% CI, 5.646-62.266; P<0.0001) levels were significantly associated with the occurrence of PD. However, no significant association was recorded for other factors. After adjusting for confounders, namely WBC count and nATP^{CD4} and sATP^{CD4} levels, patients with low sATP^{CD4} levels (<224.5 ng/ml) displayed a 15-fold higher risk of PD compared with those with high sATP^{CD4} levels (≥ 224.5 ng/ml; OR=15.392; 95% CI, 4.260-55.609; P<0.0001; Fig. 6).

Discussion

The treatment of NSCLC continues to face challenges. Notably, several multiplexed detection methods, based on an

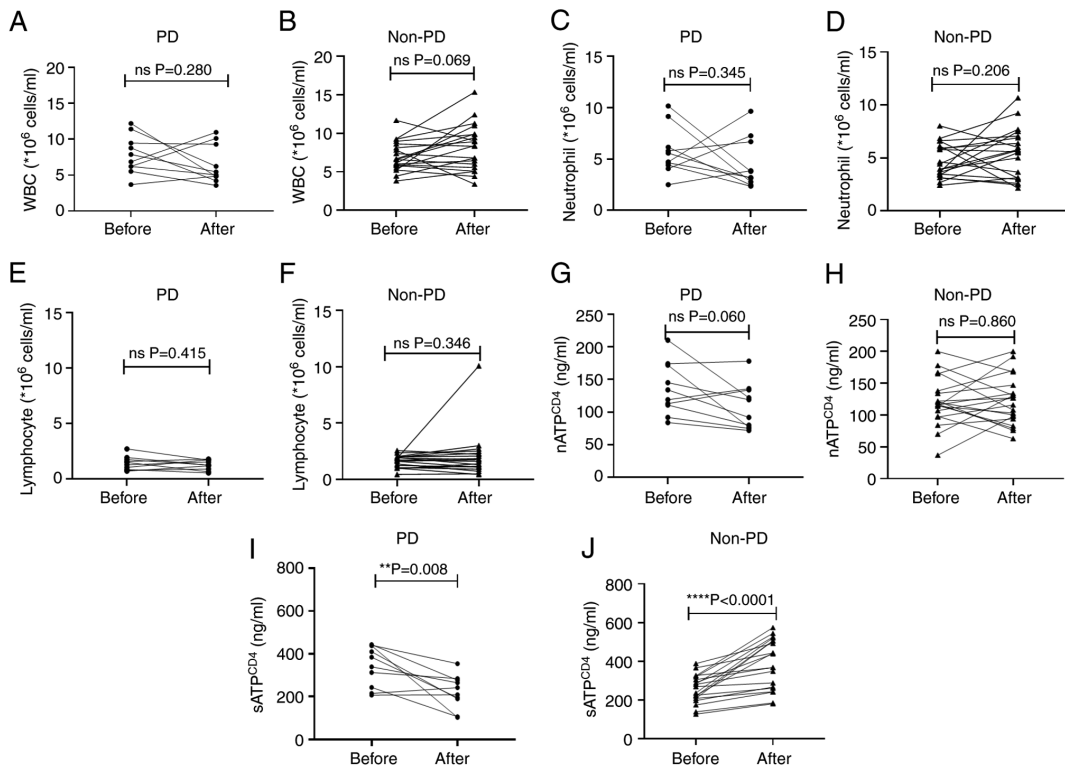


Figure 4. Changes in the peripheral blood indicators following four cycles of chemotherapy in patients in the PD (n=9) and non-PD groups (n=20). The changes in WBC in the (A) PD and (B) non-PD groups prior and after chemotherapy are shown. The changes in neutrophil count in the (C) PD and (D) non-PD groups before and after chemotherapy are presented. Changes in lymphocyte count in the (E) PD and (F) non-PD groups before and after chemotherapy. Changes in nATP^{CD4} expression prior and after chemotherapy in the (G) PD and (H) non-PD groups. Changes in sATP^{CD4} levels before and after chemotherapy in the (I) PD and (J) non-PD groups. No significant changes in WBC, neutrophil and lymphocyte count and nATP^{CD4} levels after chemotherapy were observed between the PD and non-PD groups (P>0.05). Compared with before and after chemotherapy, sATP^{CD4} levels were significantly decreased in the PD group (P=0.008), while they were notably enhanced in the non-PD group (P<0.0001) prior to chemotherapy. **P<0.01, ****P<0.0001; ns, no significant. PD, disease progression; WBC, white blood cell; nATP^{CD4}, non-stimulated CD4⁺ cell ATP concentration; sATP^{CD4}, stimulated CD4⁺T cell ATP concentration.

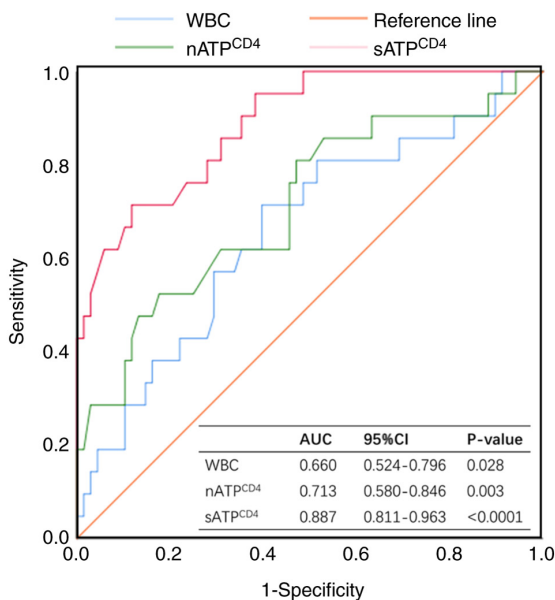


Figure 5. Receiver operating characteristics curve for diagnostic factors associated with chemotherapy efficacy in non-small cell lung cancer (n=89). The area under the curve for white blood cells was 0.660 (95% CI, 0.524-0.796), 0.713 (95% CI, 0.580-0.846) for non-stimulated CD4⁺ T cell ATP concentration and 0.887 (95% CI, 0.811-0.963) for stimulated CD4⁺ cell ATP concentration. ATP, adenosine triphosphate; CI, confidence interval; nATP^{CD4}, non-stimulated CD4⁺ cell ATP concentration; sATP^{CD4}, stimulated CD4⁺T cell ATP concentration.

optimized nanoparticle-based laser desorption/ionization mass spectrometry platform, have been developed. This treatment approach is considered as an effective tool for the screening of patients with NSCLC, potentially allowing the early diagnosis of a greater number of patients with lung cancer (25). However, for patients with advanced-stage lung cancer, chemotherapy remains a critical treatment modality. Chemotherapy efficacy is a pivotal concern in the treatment of various malignancies, since it directly affects patient outcomes and OS. However, PD remains the leading clinical problem (26,27). The current study aimed to identify factors associated with PD in patients with NSCLC undergoing chemotherapy.

The potential of immune-related biomarkers, including N6-methyladenosine and immune-related lncRNAs, in predicting immunotherapy response in lung squamous cell carcinoma via delineating molecular subtypes with varying treatment efficacy has been recently underlined (28). However, their applicability in predicting chemotherapy response has not been previously investigated. A previous study emphasized the significance of the tumor microenvironment in the context of cancer chemotherapy (29). Nevertheless, it has been suggested that the immune microenvironment status within the tumor site and the overall systemic immune profiles of patients with cancer prior to treatment can affect their response to chemotherapy (30). The association between peripheral blood-related immunological markers, including lower peripheral blood

Table II. Paired-sample area differences under the receiver operating characteristics curves for sATPCD4, nATPCD4 and WBC.

Results of the tests	z	P-value	Difference in area under the curve	Difference in standard error	95% confidence interval
sATP ^{CD4} - nATP ^{CD4}	2.592	0.010	0.174	0.326	0.042-0.306
sATP ^{CD4} - WBC	3.181	0.001	0.227	0.329	0.087-0.367
nATP ^{CD4} - WBC	0.599	0.549	0.053	0.371	-0.121-0.227

sATP^{CD4}, stimulated CD4⁺T cell ATP concentration; nATP^{CD4}, non-stimulated CD4⁺ cell ATP concentration; WBC, white blood cells.

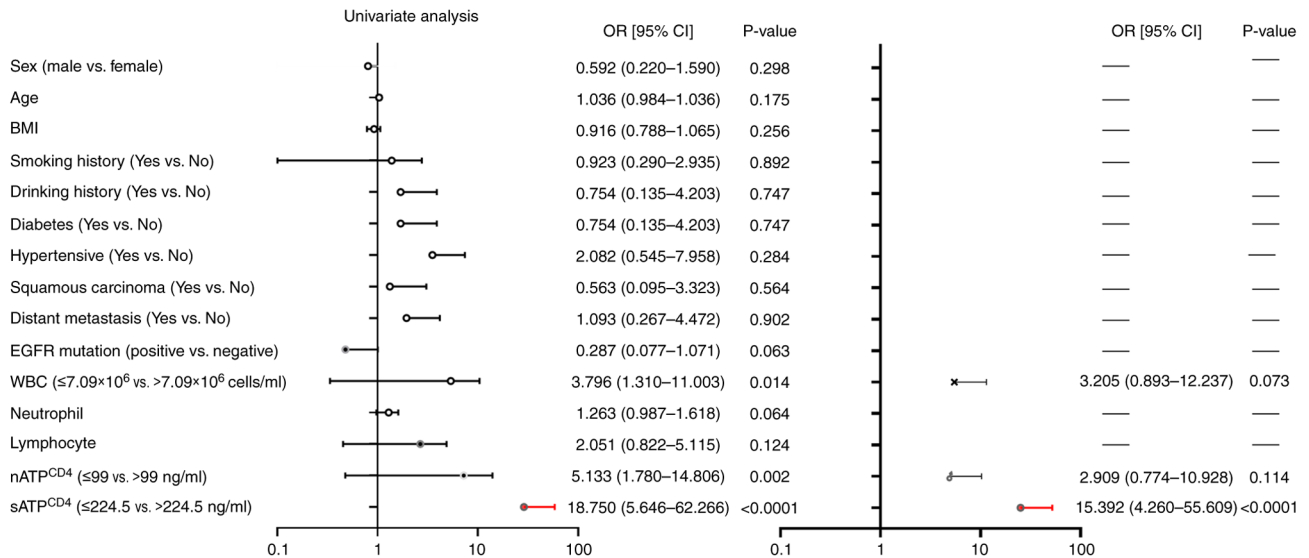


Figure 6. Univariate and multivariable analysis. Variables with $P \leq 0.05$ in univariate models were further analyzed in the multivariate analysis model. The univariate results indicated that white blood cells, non-stimulated CD4⁺ cell ATP concentration and sATP^{CD4} were risk factors for PD. After multivariable adjustment, sATP^{CD4} remained as an independent risk factor for PD. PD, disease progression; nATP^{CD4}, non-stimulated CD4⁺ cell ATP concentration; sATP^{CD4}, stimulated CD4⁺ cell ATP concentration.

mononuclear cell counts and diminished cytokine expression levels and inferior treatment responses has been widely investigated (31,32). Considering the aforementioned findings, in the present study, a comparative analysis of routine peripheral blood parameters, including WBC, neutrophil and lymphocyte counts, between the PD (n=21) and non-PD groups (n=68) was performed. The results revealed that WBC counts were notably reduced in patients in the PD group compared with those in the non-PD group, with relatively stable disease status ($P=0.028$).

The present study unveiled new possibilities for exploiting ATP detection as a sensitive and specific biomarker for predicting disease progression. The majority of cellular functions depend on ATP production and therefore intracellular ATP synthesis serves as a marker of cell activity (33). In a previous study, Cylex's Immuknow assay was used to evaluate the activity of CD4⁺T lymphocytes (34). Assessing the activity of CD4⁺T lymphocytes in the peripheral blood of organ transplant recipients can enable the early identification of potential risks associated with rejection and infection (35). Previous studies in sepsis revealed that sATP^{CD4} levels were markedly enhanced among survivors compared with non-survivors within the first day of intensive care unit admission (13,36). The aforementioned divergent findings highlight the

complex nature of sATP^{CD4} and its potential roles in several diseases. The current study revealed a noteworthy decline in sATP^{CD4} levels within the PD group following chemotherapy. Intriguingly, a notable increase in sATP^{CD4} levels was observed in the non-PD group, characterized by stable disease status (Fig. 4). Furthermore, both nATP^{CD4} and sATP^{CD4} levels were significantly diminished in patients experiencing PD compared with the non-PD group ($P=0.002$; $P<0.0001$; Fig. 3). These results supported the ability of ATP to mirror the effect of chemotherapy on immune function in NSCLC.

As a crucial factor of anti-tumor immunity, peripheral blood CD4⁺T cells play a pivotal role in regulating and enhancing the priming, migratory potential and killing activity of CTLs (37). In the present study, ROC curve analysis demonstrated a superior discriminatory ability for sATP^{CD4}, as evidenced by its higher AUC value (0.887) compared with that obtained for WBC (AUC=0.660) and nATP^{CD4} (AUC=0.713). Employing a designated cut-off value of 224.5 ng/ml for sATP^{CD4}, the sensitivity and specificity were estimated to be 77.4 and 88.2%, respectively (Fig. 5). Notably, sATP^{CD4} levels <224 ng/ml were indicative of an elevated risk of PD, thus underscoring a robust association between CD4⁺T cell immune function and patient prognosis. Low ATP levels indicate a state of immunosuppression, thus accelerating tumor

progression via the intricate immunosuppressive network established by interactions between cancer cells and host immune cells. This phenomenon could promote tumor growth, while simultaneously providing a shield against immune attacks, thus contributing to the complex dynamics of cancer progression (38). It has been reported that during the complex immune editing process, where several CD4⁺ T cell subsets play significant roles in both tumor promotion and rejection, sATP^{CD4} serves as a valuable indicator reflecting the overall net status of immune activity (35).

Factors that affect treatment efficacy in patients with NSCLC include lymphocyte subpopulations (39,40), treatment regimen (41,42) and gene mutations (42,43). The results of the present study showed that only sATP^{CD4} was a risk factor for tumor progression. This finding could be associated with the small sample size. The results of the current study were consistent with those reported in liver cancer, demonstrating that patients with low sATP^{CD4} levels exhibited markedly lower PFS and OS compared with those in the high ATP group (18). Another study also suggested that adjusting the dosage of immunosuppressive agents based on ATP levels could markedly improve the OS and reduce infection rates in liver transplant patients (44). The present study and previous studies demonstrated that sATP^{CD4} levels could reflect immune function in patients with cancer and were associated with treatment efficacy and prognosis. However, the mechanism underlying the effect of sATP^{CD4} levels on reflecting the efficacy of chemotherapy remains to be elucidated.

In the present study, the particular underlying mechanism was investigated, which could be associated with how chemotherapy could promote the death of tumor cells within a short period. In turn, the death of tumor cells induces the release of several tumor antigens, potentially stimulating antigen-presenting cells to present tumor antigens to T cells, thus promoting T cell activation and proliferation (20). Additionally, the combination of EGFR-targeted agents could enhance the presentation of MHC I-class tumor antigens to facilitate the uptake of tumor material by dendritic cells and promote T cell activation in the absence of additional immune stimulation signals (45). Activated T cells infiltrate tumor tissues and attack antigen-expressing tumor cells. Furthermore, emerging evidence has suggested that cellular metabolism plays a critical role in T cell differentiation and function. Changes in the metabolic activity of T cells can directly affect their function and survival, which is reflected in ATP expression levels (46). Therefore, higher nATP^{CD4} and sATP^{CD4} expression levels were observed in patients with disease stabilization or remission following chemotherapy. Conversely, in patients with PD, chemotherapy drugs could not only promote tumor cell injury, but they could also harm normal immune cells, thus resulting in impaired immune function. The aforementioned processes could be accompanied by lower nATP^{CD4} and sATP^{CD4} expression levels. Consequently, the immune cells of these patients could fail to effectively clear tumor cells, thus leading to PD.

However, the present study had several limitations. First, the relatively small sample size and limited number of events, as well as the fact that it was conducted only on Chinese individuals without validation in other ethnic backgrounds, constrained comprehensive multivariable analysis. Additionally, an increasing number of studies have

demonstrated that advanced NSCLC patients with KRAS mutations exhibit markedly lower objective response rate and potentially lower 6-month/1-year PFS rates compared with wild-type patients after first-line chemotherapy (43,47,48). However, the limited number of cases with detected KRAS mutations in this study (Table I) precludes the calculation of whether KRAS gene mutations affect chemotherapy efficacy. Whether KRAS mutations serve as risk factors for sATP^{CD4} requires further investigation. Therefore, further validation of the results is needed to draw definitive conclusions. Additionally, the retrospective and single-center nature of this study may introduce some bias, potentially resulting in inherent deviations. Considering the primary goals of immunotherapy, extending the observation period, treatment cycles and longitudinally monitoring changes in sATP^{CD4} subsets are crucial for investigating their association with OS and PFS.

In summary, in the present study a noteworthy association between low nATP^{CD4} and sATP^{CD4} levels and tumor progression was observed in patients with advanced NSCLC treated with chemotherapy. These findings could assist physicians in assessing immune function to clearly determine the immunological status of patients with cancer, thus tailoring treatment strategies to prevent disease progression. The current study was the first to provide such findings, to the best of the authors' knowledge. However, further prospective and multicenter studies with larger sample sizes are needed to validate the aforementioned results.

The present study suggested that sATP^{CD4} levels could serve as an indicator of the effect of chemotherapy on immune function and holds promise for assessing the potential risk of disease progression in patients with NSCLC.

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Availability of data and materials

The data generated in the present study may be requested from the corresponding author.

Authors' contributions

WY, KH, NT, WL, ZT, QH, DY, HL, ZD, YX and GY contributed to the study conception and design. Material preparation, data collection and analysis were performed by WY, KH and NT under supervision of GY. The first draft of the manuscript was written by WY. KH and NT confirm the authenticity of all the raw data. WL, ZT and QH reviewed and revised the first draft, conducted domestic and international literature searches for the discussion section and improved the discussion section.

DY and HL further reviewed and examined the data analysis and graphics production in the article. ZD and YX formatted the final version of the article. GY supervised the entire experiment and performed the final review of the article. All authors read and approved the final manuscript.

Ethics approval and consent to participate

The present study received approval from the Ethics Committee of the Fifth Affiliated Hospital of Guangzhou Medical University (Guangzhou, China; dated 08-07-2022; approval no. KY01-2022-07-08). Written informed consent was obtained from all participants, allowing the use of their clinical data in this study. The research adhered to the principles outlined in The Declaration of Helsinki.

Patient consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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