



Neoadjuvant camrelizumab, nab-paclitaxel, and carboplatin in patients with stage IB-IIIa non-small cell lung cancer (NANE-LC): a study protocol of prospective, single-arm, multicenter, phase II study

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Background: Previous studies have shown that neoadjuvant immune checkpoint inhibitors (ICIs) combined with chemotherapy in patients with stage IB–IIIa non-small cell lung cancer (NSCLC) significantly improved the major pathological response (MPR) and the pathological complete response (pCR) rates. However, high-level evidence-based medical data confirming this effect are still lacking. In addition, there is an urgent need to develop an appropriate strategy to predict the benefit for patients receiving ICIs. In this study, we describe an ongoing study on the effect of neoadjuvant therapy with camrelizumab, nab-paclitaxel, and carboplatin on stage IB–IIIa NSCLC patients. The aim of this study is to establish a multiomics artificial intelligence system for predicting neoadjuvant therapy efficacy and assisting decision-making.

Methods: This prospective, single-arm, multicenter, phase II trial will enroll a total of 40 patients who will undergo surgery after three cycles of neoadjuvant therapy with camrelizumab, nab-paclitaxel, and carboplatin. The MPR rate is the primary endpoint, while the rates of pCR, complete resection, objective response, disease-free survival (DFS), adverse events (AEs), and quality of life (QOL) are secondary endpoints. Exploratory endpoints will serve to establish a multiomics artificial intelligence system for neoadjuvant therapy effect prediction and decision-making assistance based on radiomics, metabolism, genetic, and clinic-pathological characteristics and to explore the mechanisms of drug resistance.

Discussion: The efficacy of ICIs is influenced by many factors, including patient's driver genes and smoking status. Thus, further subgroup analysis is needed. This study will indicate if our new multiomics artificial intelligence system constitutes a valid strategy for neoadjuvant therapy effect prediction and decision-making assistance in the context of neoadjuvant camrelizumab, nab-paclitaxel, and carboplatin treatment for patients with stage IB–IIIa NSCLC.

Trial Registration: This trial has been registered at ClinicalTrials.gov (identification number: NCT04541251).

Keywords: Neoadjuvant therapy; non-small cell lung cancer (NSCLC); immune; checkpoint inhibitors; artificial intelligence; major pathological response

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Introduction

Randomized controlled trials have shown that patients with non-small cell lung cancer (NSCLC) who receive neoadjuvant chemotherapy obtain enhanced survival benefits compared with patients treated with surgery alone (1-3). In the NATCH trial, NSCLC patients achieved 22.8% of major pathological response (MPR), and virtually no patients achieved pathological complete response (pCR) after neoadjuvant chemotherapy (2). Although neoadjuvant chemotherapy improves the survival outcome of resectable NSCLC, it is still insufficient, and new neoadjuvant therapy regimens are needed to improve treatment efficacy.

Immune checkpoint inhibitors (ICIs) that target the programmed cell death 1 (PD-1) axis have revolutionized the management of advanced NSCLC (4-8). These ICIs have been utilized for neoadjuvant therapy in patients with resectable NSCLC. The patients with resectable NSCLC who were enrolled in the CheckMate 159 trial (9) were treated with two cycles of neoadjuvant therapy using nivolumab monotherapy. This treatment significantly improved the MPR rate (45%) and the pCR rate (15%). However, the NEOSTAR (10) and the LCMC3 (11) trials, assessing patients under neoadjuvant ICI monotherapy, yielded an MPR rate of 17% and 19%, respectively. Thus, the benefit of neoadjuvant therapy based on ICI monotherapy in patients with resectable NSCLC still remains controversial.

To further improve the benefits for patients with resectable NSCLC, a new strategy of neoadjuvant therapy combining ICIs and chemotherapy has been implemented. The NADIM trial (12) was a phase II clinical trial that enrolled stage IIIA NSCLC patients who received three cycles of neoadjuvant therapy with nivolumab, paclitaxel, and carboplatin. This trial demonstrated that neoadjuvant ICIs combined with chemotherapy significantly improve patients' MPR and pCR ratios (83% and 71%, respectively). Another phase II clinical trial (13) enrolled patients with stage IB-III A NSCLC who received four cycles of neoadjuvant therapy with atezolizumab, nab-paclitaxel, and carboplatin. In this trial, 50% of the patients achieved MPR, of which 21% achieved pCR. Although both trials

demonstrated that neoadjuvant ICIs combined with platinum-based chemotherapy bring additional benefits for patients with NSCLC, high-level evidence-based medical data are still lacking to support this observation.

Moreover, current published studies have focused on finding appropriate biomarkers or strategies to predict the benefit for patients receiving ICIs. Among these, the expression of programmed cell death-ligand 1 (PD-L1), the tumor mutation burden (TMB), and high microsatellite instability or mismatch repair deficiency have been widely used for patients with advanced solid tumors (6,14-16). The combination of these biomarkers with other factors, such as sex, genetic mutations, or ctDNA maximum somatic allele frequency, seems to be a promising strategy for predicting the benefit for patients receiving ICIs (17-20). Regarding patients with resectable NSCLC, the Checkmate-159 trial indicated an absence of correlation between pathological response to ICIs and PD-L1 expression, while there was a linear correlation between pathological response and TMB (9).

Currently, some new strategies for predicting the efficacy of ICIs are being discovered and reported. The INSPIRE trial (21), a phase II trial, demonstrated that the changes in ctDNA before and after ICIs are strongly correlated with the benefit of ICIs in patients with incurable, locally advanced, or metastatic solid tumors. Importantly, the CheckMate 078 and CheckMate 870 (22) trials revealed a strong association between the gut microbiome diversity and patient's response to ICIs, implying that gut microbiome diversity may be a predictive strategy for ICIs in patients with NSCLC. Furthermore, long non-coding RNAs (lncRNAs) were found to be closely related to immune dysfunction and response to ICIs. A novel multiomics algorithm model that included TMB, PD-L1 expression, cytotoxic T-lymphocyte infiltration, and lncRNA score displayed a strong correlation with the overall survival (OS) of patients treated with ICIs (23). Moreover, some studies have shown that artificial intelligence can be applied for the diagnosis and treatment of NSCLC patients, while radiographic characteristics might serve as noninvasive approaches for predicting the response to immunotherapy (24,25). However, it is still unclear whether these predictive

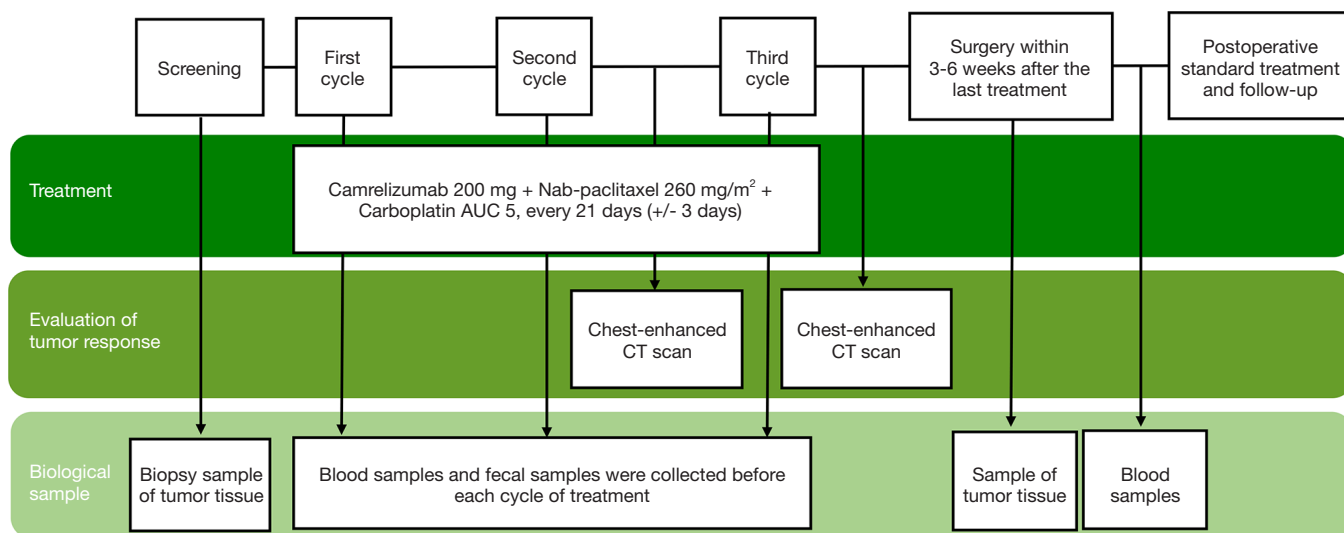


Figure 1 Flowchart of the trial. AUC, area under the curve; CT, computed tomography.

biomarkers or strategies are effective in predicting the benefit to patients receiving neoadjuvant ICIs.

The aim of this study is to evaluate the efficacy and safety of neoadjuvant therapy with the PD-1 inhibitor camrelizumab, nab-paclitaxel, and carboplatin for patients with stage IB-III A NSCLC and to establish a multiomics artificial intelligence system to predict neoadjuvant therapy efficacy based on radiomics, metabolism, genetic, and clinical-pathological characteristics. In addition, the resistance mechanisms to neoadjuvant therapy that utilizes camrelizumab, nab-paclitaxel, and carboplatin will be explored.

Methods

Study design

This NANE-LC study is a prospective, single-arm, multicenter, phase II trial. A total of 40 patients will undergo surgery after three cycles of neoadjuvant therapy with camrelizumab, nab-paclitaxel, and carboplatin. *Figure 1* shows the flowchart of the trial.

Eligibility criteria

The key inclusion and exclusion criteria of this study are shown in *Figure 2*.

Treatment

The patients will receive three cycles (one cycle is defined

as a period of 21 ± 3 days) of neoadjuvant therapy with camrelizumab 200 mg, nab-paclitaxel 260 mg/m², and carboplatin area under the curve (AUC) 5. This treatment will be followed by surgery.

Patient registration

After confirming eligibility and obtaining a signed informed consent form, each patient will be registered and will receive treatment. Patient recruitment began in August 2020. It is expected to continue for the next two years.

Specimen collection

During the trial, samples of tumor tissues will be collected, including biopsy samples and surgical tumor tissue samples. Blood samples and fecal samples will be collected before every cycle of neoadjuvant therapy and before surgery. The tumor tissues and blood samples will be used for whole genome sequencing, whole transcriptome sequencing, and single-cell sequencing. The fecal samples will be used for 16S rDNA sequencing and nontargeted metabolomics.

Dosages and treatment regimen

Adjustment of the dosage regimens in case of adverse events (AEs) occurring during the trial shall be based on the dose-adjustment regulations specified on the drug medication guides or relevant clinical decisions taken by the

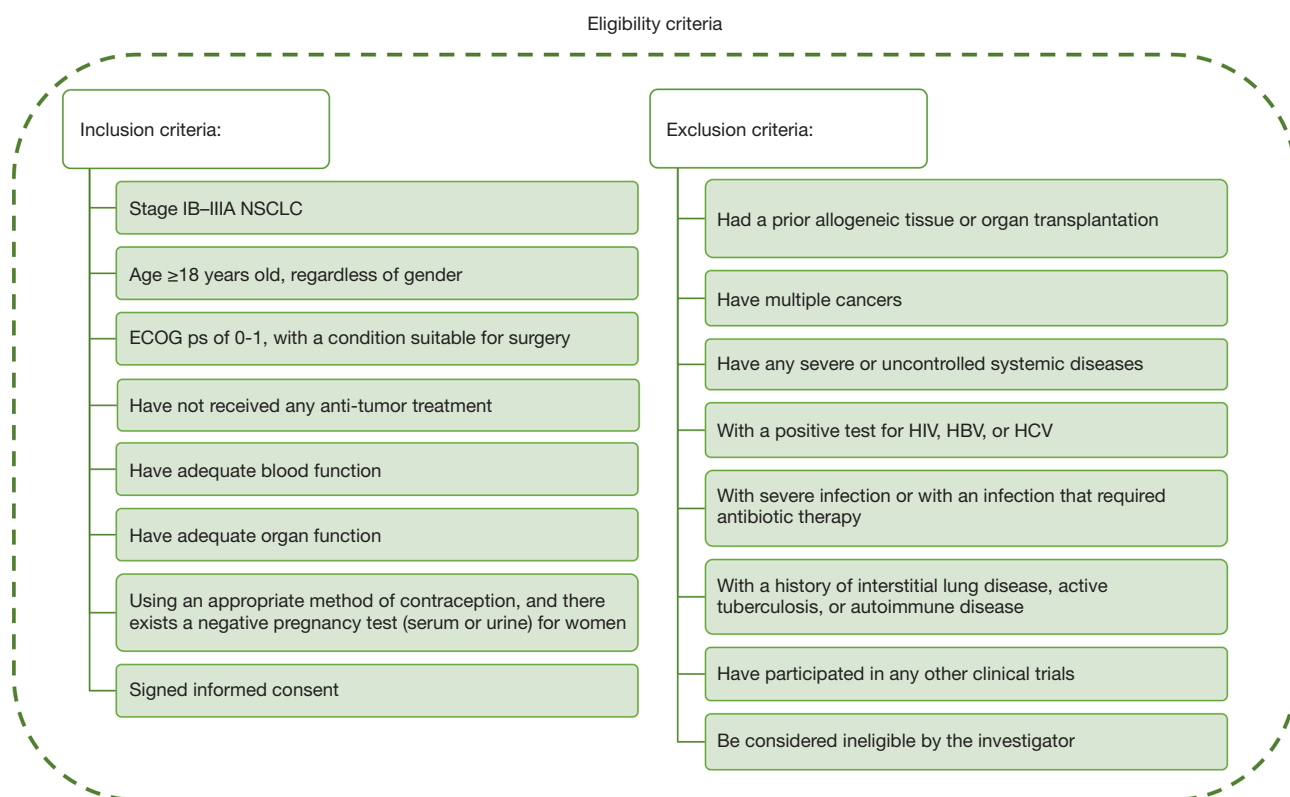


Figure 2 Key inclusion and exclusion criteria. NSCLC, non-small cell lung cancer; ECOG PS, Eastern Cooperative Oncology Group performance status; HIV, human immunodeficiency virus; HBV, hepatitis B virus; HCV, hepatitis C virus.

researchers.

Endpoint measures

The primary endpoint measure is the MPR rate. The secondary endpoint measures are the pCR rate, the complete resection (R0) rate, the disease-free survival (DFS), the OS, the objective response rate (ORR), AEs, serious AEs, and quality of life (QOL). The exploratory endpoint measure is the establishment of an artificial intelligence prediction system for neoadjuvant therapy effect prediction and exploration of drug resistance mechanisms. This system will be based on the multiomics data, including radiomics, metabolism, genetic, and pathological characteristics. A detailed description is provided in *Figure 3*.

Response and safety evaluation

Pre-treatment evaluations of the patients are compulsory and include chest and abdomen-enhanced computed tomography (CT) scans, brain magnetic resonance imaging

scans, bone scans or positron emission tomography scans, and electrocardiography. Patients will undergo a tumor evaluation using a chest-enhanced CT scan before the third cycle of neoadjuvant therapy and prior to surgery. The tumor response and/or radiologic disease progression will be evaluated based on version 1.1 of the Response Evaluation Criteria in Solid Tumors. AEs will be recorded using version 5.0 of the National Cancer Institute's Common Terminology Criteria for Adverse Events. The QOL will be assessed using questionnaires of the Quality of Life Questionnaire-Core 30 of The European Organization for Research and Functional Assessment of Cancer Therapy-Lung before and after every cycle of neoadjuvant treatment.

Statistical analysis

Descriptive statistics will be used to summarize the demographic data and baseline characteristics of the patients, and a *t*-test, a chi-square test, or a rank sum test will be used to assess group equilibria. The number and

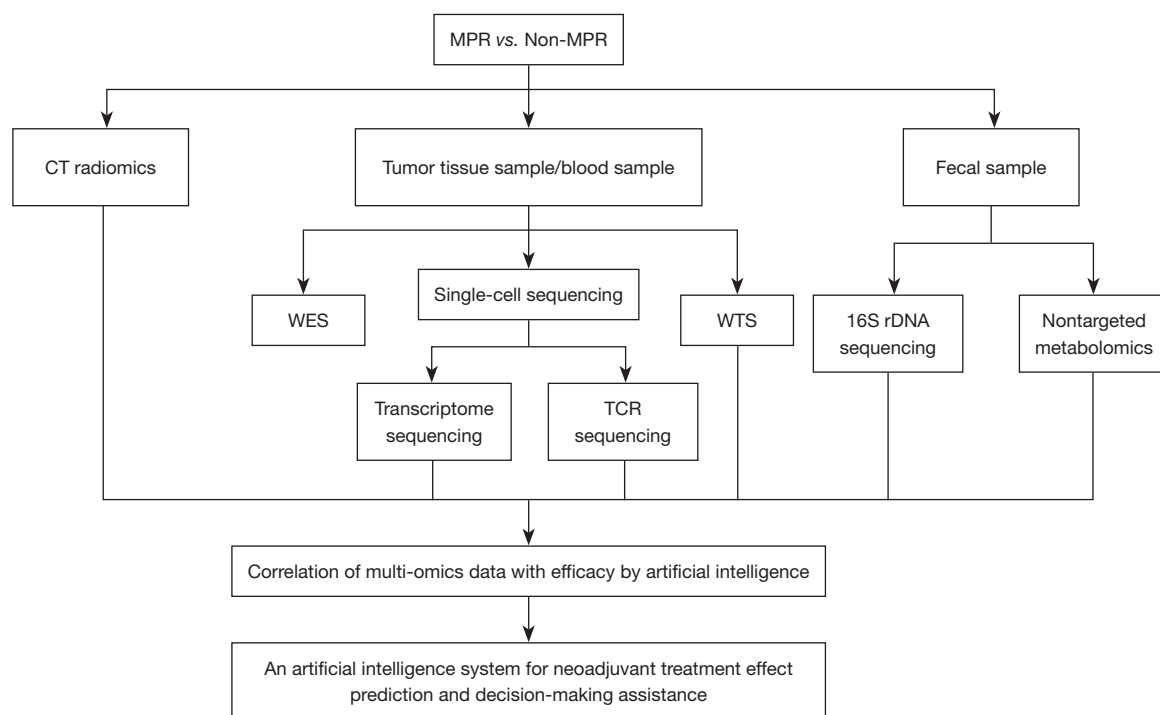


Figure 3 Detailed description of the artificial intelligence prediction system used to assess neoadjuvant-based therapy efficacy. MPR, major pathologic response; CT, computed tomography; WES, whole genome sequencing; WTS, whole transcriptome resequencing; TCR, T cell receptor.

percentage of MPR, pCR, the R0 excision rate, and the ORR will be calculated. The Kaplan-Meier method will be used to estimate the survival curve, median, and annual rate of DFS and OS. For the treatment sensitivity analysis of the biomarkers, the Cox univariate and multivariate analyses will be used to analyze the correlation between biomarker expression, clinical efficacy, and prognosis. Artificial intelligence algorithms will be used to establish a prediction system for the neoadjuvant therapy outcome based on the multiomics data. In the process of Artificial intelligence analysis, convolutional neural network (CNN)-based deep learning will be used to train images, and a variety of algorithms, including GoogLeNet, ResNet, RCNN, fast-rCNN, fath-rCNN, and VGGNet, will be used for feature extraction, screening, modeling, and verification. Moreover, transfer learning algorithm will be used, which refers to the method of transferring the pre-trained CNN model to other datasets and relearning the characteristics of the target dataset. Data augmentation will be adopted to increase the size of the training dataset and expand the existing samples by iterations of random translational shift, rotation, and horizontal and vertical

flips. The performance of the prediction system will be evaluated using the receiver operating characteristic curve, the corresponding AUC, the diagnostic accuracy, sensitivity, specificity, the positive predictive value, and the negative predictive value. The Clopper Pearson method will be used to calculate the 95% confidence interval. All statistical analyses will be performed with R software 4.0.5 and SPSS 24.0 software.

Sample size

The sample size was calculated using the Simon two-stage method. MPR rates of 22.8% and 17% were expected according to the NATCH (2) and the NEOSTAR (10) trials, respectively, and an MPR rate of 19% was expected according to the LCMC3 trial (11), in which patients received neoadjuvant ICI monotherapy. It is believed that patients treated with neoadjuvant ICI monotherapy could achieve MPR rates of about 20%. Therefore, an MPR rate of 42.8% was expected in this study, with a two-sided alpha value of 0.05 and a power of 0.8. The sample size for the first stage was 13. If more than three cases reach the

MPR, the second stage will begin, in which 27 cases will be included. The results of this trial will be considered positive if more than 13 cases out of 40 effective cases reach MPR. Effective cases are defined as patients who received at least once cycle of neoadjuvant therapy and completed the first tumor assessment.

Ethical consideration and registration

This trial protocol was approved on July 23, 2020 by the ethics committee of Sun Yat-sen Memorial Hospital of Sun Yat-sen University (No. 2020-KY-061-001), Jiangxi Provincial Cancer Hospital (No. 2021ky074), and Third Affiliated Hospital of Sun Yat-sen University (No. [2021]02-205-01). All patients enrolled in the trial have signed or will sign informed consent forms prior to registration according to the Declaration of Helsinki (as revised in 2013). This protocol has been registered at ClinicalTrials.gov (identification number: NCT04541251). As of August 2021, three subjects have been enrolled and have signed the informed consent forms.

Discussion

ICIs, one of the most significant immunotherapies, have entered clinical practice and brought new hope to NSCLC patients. Although many ongoing trials focus on the identification of new biomarkers, such as PD-L1 and TMB, usable as potential surrogates for the prediction of clinical outcomes, current strategies and biomarkers for predicting ICI efficacy are still limited. In this study, we not only aim to evaluate the efficacy and safety of ICIs combined with chemotherapy as neoadjuvant therapy for NSCLC patients but also to explore the efficacy of new prediction strategies.

It is well known that NSCLC patients with epidermal growth factor receptor (EGFR) mutations are preferred for targeted therapy and benefit little from immunotherapy. Related studies have shown that ICIs combined with chemotherapy or target drug therapy might show improved efficacy for NSCLC patients with EGFR mutations (26-28). NSCLC patients with unknown EGFR status could benefit from ICIs alone (9) or combined with chemotherapy (11,13). Therefore, we did not use a particular EGFR status in the enrollment criteria, and we hope to discuss this point during the future data analysis.

In the meantime, a significant correlation between PD-L1 expression and smoking habits has been reported (29). Yet, recent subgroup analyses from the KEYNOTE-042

study (30) showed that there was no significant OS benefit for the never-smoking subgroup of patients with PD-L1 positive expression, regardless of PD-L1 TPS $\geq 1\%$, $\geq 20\%$, or $\geq 50\%$. In contrast, subgroup analyses in the KEYNOTE-189 study (4) showed better OS benefit for never-smoking NSCLC patients than for current or former smoking patients who received ICI combined with chemotherapy. In a recent retrospective study (31), the researchers analyzed the association between smoking and ICI activity in 315 NSCLC patients with PD-L1 TPS $\geq 50\%$. They found that a number of never smokers and light smokers significantly benefited from immunotherapy. Although higher TMB levels in heavy smokers might lead to more significant benefits from immunotherapy, smoking status alone did not fully explain the complexity of neoantigen production and presentation. The influence of patients' smoking habits on ICIs' efficacy is worthy of further subgroup analysis.

With the smooth completion of this study, we hope to deliver new evidence on patients with stage IB-IIIa NSCLC who received neoadjuvant camrelizumab, nab-paclitaxel, and carboplatin. Based on radiomics, metabolism, genetic, and clinical-pathological characteristics, our new multiomics artificial intelligence system may represent a new valuable strategy for neoadjuvant therapy efficacy prediction and decision-making. In addition, this new tool may provide potential directions for exploring the mechanisms of resistance to current cancer therapies.

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Footnote

Data Sharing Statement: Available at <https://dx.doi.org/10.21037/jtd-21-1022>

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Conflicts of Interest: All authors have completed the ICMJE uniform disclosure form (available at <https://dx.doi.org/10.21037/jtd-21-1022>). The authors have no conflicts of interest to declare.

Ethical Statement: The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved. This trial protocol was approved on July 23, 2020. It was supervised and managed by the ethics committee of Sun Yat-sen Memorial Hospital of Sun Yat-sen University (No. 2020-KY-061-001), Jiangxi Provincial Cancer Hospital (No.2021ky074), and Third Affiliated Hospital of Sun Yat-sen University (No. [2021]02-205-01). All patients enrolled in the trial have signed or will sign informed consent forms prior to registration, in accordance with the Declaration of Helsinki (as revised in 2013). As of August 2021, three subjects have been enrolled and have signed the informed consent form.

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