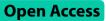
RESEARCH

BMC Pulmonary Medicine



Impact of immune checkpoint inhibitors (ICIs) therapy on interferon-γ release assay (IGRA) and diagnostic value in non-small cell lung cancer (NSCLC) patients

Yijiao Xu^{1,2†}, Qingwei Zhang^{1,2†}, Zhisheng Chen^{1,2†}, Shuwen Yang^{1,2}, Haiyan Chen^{1,2}, Xiong Xiao^{1,2*} and Hongni Jiang^{1,2,3,4*}

Abstract

Background Tuberculosis (TB), a highly contagious respiratory disease, presents a significant global health threat, with a notable increase in incidence reported by the WHO in 2022. Particularly, the interplay between TB and non-small cell lung cancer (NSCLC) gains attention, especially considering the rising use of immune checkpoint inhibitors (ICIs) in cancer treatment. This interplay may influence TB diagnostics and reactivation, warranting a closer examination.

Methods A retrospective analysis was conducted on clinical data of NSCLC patients with positive T-SPOT results before undergoing anti-tumor treatment at Zhongshan Hospital (Xiamen), Fudan University, from January 1, 2021 to December 31, 2022. We assessed the incidence of tuberculosis reactivation and treatment outcomes among these patients. Moreover, we compared the differences in tuberculosis activity between the ICIs and non-ICIs treatment groups. Additionally, we observed the changes in T-SPOT spot count before and after immunotherapy, analyzing their association with tuberculosis activity and prognosis.

Results A total of 40 NSCLC patients with positive T-SPOT results before treatment were included in the study, with 26 patients in the ICIs treatment group and 14 patients in the non-ICIs treatment group. The study found no significant differences between the two groups in terms of gender, age, stage, histological type, performance status, driver gene expression, and distant metastasis. With a median follow-up time of 10.0 (6.0-14.5) months, three cases (11.5%) in the ICIs treatment group developed tuberculosis activity, diagnosed at 2, 3, and 12 months after ICIs treatment initiation. Conversely, no tuberculosis activity was observed in the non-ICIs treatment group, and the difference between the two groups was not significant (P=0.186). Among the 32 patients who received

[†]Yijiao Xu, Qingwei Zhang, Zhisheng Chen authors contributed equally to this work.

*Correspondence: Xiong Xiao xiao.xiong@zsxmhospital.com Hongni Jiang jiang.hongni@zs-hospital.sh.cn

Full list of author information is available at the end of the article



© The Author(s) 2024. **Open Access** This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit http://creativecommons.org/licenses/by/4.0/. The Creative Commons Public Domain Dedication waiver (http://creativecommons.org/publicdomain/zero/1.0/) applies to the data made available in this article, unless otherwise stated in a credit line to the data.

ICIs treatment, spot count dynamics were diverse: four cases (12.5%) showed an increase, 12 cases (37.5%) had no change, and 16 cases (50.0%) had a decrease. During the follow-up, the progression rate (PD) was 50.0%, 75.0%, and 62.5% in the three groups, respectively (P=0.527). Similarly, the mortality rate was 0%, 25.0%, and 25.0%, respectively (P=0.106). Interestingly, among the patients with decreased spot counts, three cases (18.75%) were diagnosed with active pulmonary tuberculosis.

Conclusions For NSCLC patients with a positive T-SPOT response undergoing ICIs treatment, our study observed indications of active tuberculosis. The varied T-SPOT spot count changes post-ICIs treatment suggest a complex interaction, potentially linking T-SPOT spot count reduction to tuberculosis reactivation risk. These preliminary findings underscore the importance of further research to more accurately assess T-SPOT's diagnostic utility in this context.

Keywords Non-small cell lung cancer (NSCLC), Immune checkpoint inhibitors (ICIs), T-SPOT, Tuberculosis, Reactivation, Treatment outcomes

Background

Tuberculosis (TB) is a highly contagious respiratory disease that presents a grave global threat. The 2022 report from the World Health Organization (WHO) reveals an unsettling change in the TB incidence trend, which had been declining over the past two decades. Contrary to this previous trend, there was an alarming increase of 3.6% in TB incidence from 2020 to 2021, affecting approximately 10.6 million individuals worldwide [1]. Research has shown that individuals with a history of TB are at a significantly higher risk of developing lung cancer. In fact, the incidence of lung cancer is increased by 3-10 times in patients with a previous TB infection compared to those without [2, 3]. In lung cancer patients, TB can manifest either as a concurrent or a latent infection, complicating the treatment of the tumor [4, 5]. The Interferon-Gamma Release Assay (IGRA) is one of the diagnostic methods used for detecting tuberculosis infection and is recommended by the WHO as a primary in vitro immunological test for diagnosing Mycobacterium tuberculosis(MTB, commonly known as TB) infection, with T-SPOT.TB being one of the most commonly used versions. This assay functions by detecting the production of IFN-y by effector T cells in the peripheral blood of the subject following stimulation with MTB-specific antigens. The enzyme-linked immune spot assay (ELISPOT) is used to quantify the number of IFN-y-releasing effector T cells, which determines the test result [6].

In recent years, immune checkpoint inhibitors (ICIs) have been increasingly used in the treatment of various malignancies, including lung cancer and melanoma [7, 8]. ICIs have the potential to release the suppression of T cells in the human body, enhancing the anti-tumor activity of the immune cells. This mechanism could also enhance the anti-mycobacterial activity against tuberculosis. However, previous literature has shown an increased incidence of active TB or conversion to positive results in interferon-gamma release assays (IGRAs) during ICIs treatment [9]. The prolonged use of ICIs in NSCLC patients can lead to T cell activation and

exhaustion, reducing T cell numbers and potentially increasing susceptibility to T cell-dependent infectious diseases [10, 11]. Given that T-SPOT results are also dependent on T cells. Thus, ICIs therapy in lung cancer patients could potentially affect T-SPOT outcomes, thereby impacting the diagnosis of pulmonary tuberculosis.

To date, the relationship between ICIs therapy in lung cancer and T-SPOT outcomes remains underexplored. A solitary prospective study reported a 3.3% incidence of T-SPOT conversion (from negative to positive) in lung cancer patients undergoing ICIs therapy, with 1.6% developing active pulmonary tuberculosis. However, this study included only 18 cases with positive T-SPOT results before treatment [12] without delving into the relationship between T-SPOT outcomes, immunotherapy efficacy, and prognosis.

Consequently, this research focuses on a T-SPOT.TB study aiming to monitor T-SPOT before and after ICIs therapy to examine the impact of ICIs treatment on the outcomes of the interferon-gamma release assay and its diagnostic efficacy in NSCLC patients.

Methods

Study design and study approval

The medical records of patients with non-small cell lung cancer (NSCLC) who received treatment at Zhongshan Hospital (Xiamen) Fudan University between January 1, 2021 and December 31, 2022 were retrospectively reviewed. Data were collected from our electronic medical record and patient follow-up. The data collected for all patients included age, sex, performance status (PS), tumor-node-metastasis (TNM) staging, histology, treatment regimens, survival, and adverse events.

Inclusion Criteria:1) Age greater than 18 years. 2) Diagnosis of stage IV NSCLC or stage IIIB/IIIC NSCLC that was either unresectable or had postoperative recurrence, confirmed both pathologically and radiologically. 3) Completion of a minimum of 2 cycles of anti-tumor treatment, which could include immune checkpoint inhibitors (ICIs), chemotherapy, or combination therapy. 4) Completion of T-SPOT testing prior to the commencement of treatment. 5) Availability of complete and standardized treatment and follow-up data. Exclusion Criteria:1) Diagnosis of small cell lung cancer (SCLC). 2) Presence of severe comorbidities affecting vital organs such as the heart, brain, liver, or kidneys. 3) Diagnosis with active pulmonary tuberculosis or extrapulmonary tuberculosis prior to treatment. 4) Required long-term (defined as continuous use for more than 4 weeks) oral or intravenous administration of corticosteroids. 5) Current treatment with other immunosuppressive agents.

This study was approved by the ethics committee of Zhongshan Hospital, Xiamen Branch, Fudan University (No. B2022-068).

Evaluation of responses

We evaluated the overall survival (OS) and progressionfree survival (PFS) among patients. OS was defined as the duration from the commencement of immunotherapy until death from any cause. Similarly, PFS was determined as the interval from the initial dose of immunotherapy to either the documented progression of the disease or death, whichever occurred first. The assessment of treatment efficacy was conducted according to the Response Evaluation Criteria in Solid Tumors (RECIST) version 1.1 [13]. Data analysis was conducted up to the censored date of May 31, 2023, with a median follow-up duration of 13.0 (interquartile range: 9.5–18.5) months.

Assessment of T-SPOT.TB

In this study, the T-SPOT.TB assay (Oxford Immunotec Ltd., UK) was utilized. Initially, fresh venous blood samples were collected from the patients, and peripheral blood mononuclear cells (PBMCs) were isolated. These cells were then resuspended in AIM-V medium to achieve a concentration of 2.5×10^5 cells/mL. For the assay, 50 µL of this cell suspension was dispensed into each of the four wells of an IFN- γ antibody-coated microplate. AIM-V medium acted as the negative control, while PHA served as the positive control. ESAT-6 and CFP-10 were added to test wells A and B, respectively. The microplate was then incubated at 37 °C in a CO2 incubator for 16-20 h. After incubation, the microplate was washed, and an alkaline phosphatase-conjugated secondary antibody was applied. Subsequently, the plate underwent another incubation at 4 °C for 1 h and was washed with PBS. To visualize the results, BCIP/NBT substrate solution was added, inducing a color change. The spots formed were counted and interpreted using a microscope. A test was considered positive if: (1) the net number of spots in the test well (after subtracting the count from the negative control well) was ≥ 6 , provided the count in the negative control well was between 0 and 5, or (2) the count in the test well was at least double that in the negative control well, assuming the negative control had ≥ 6 spots.

Acid-fast staining of sputum smears

Sputum specimens were collected from patients in the early morning, with a focus on collecting purulent sputum. When required, specimens were diluted with normal saline and preserved in a sterile, sealed container. For analysis, a small aliquot of the specimen was carefully placed on a glass slide and subjected to the acidfast staining technique. Following staining, the slide was meticulously examined under a microscope, with 100 fields of view being scrutinized. A result was deemed positive if 3 or more acid-fast bacilli were identified within any single field of view.

Statistical analysis

For the analysis of continuous data that followed a normal distribution, the mean and standard deviation were represented as $(X\pm SD)$. In contrast, for continuous data not adhering to a normal distribution, the median and interquartile range (IQR) served as the descriptive statistics. Categorical data were summarized using frequencies and percentages. To compare continuous data across groups, t-tests or non-parametric tests were employed based on the data distribution. For categorical data comparisons, the chi-square test was the primary method used. In instances where the expected frequency in any table cell was below 5, Fisher's exact test was utilized instead. A p-value of less than 0.05 was deemed to indicate statistical significance. All statistical analyses were conducted using SPSS software, version 26.0.

Results

Impact of ICIs therapy on tuberculosis activity in T-SPOTpositive NSCLC patients

Patient characteristics

In this study, a cohort of 40 T-SPOT-positive non-small cell lung cancer (NSCLC) patients was analyzed before the initiation of any treatment (Fig. 1). The gender distribution was predominantly male with 32 (80.0%) patients, while females accounted for 8 (20.0%). The median age of the participants was 65 years, with a division of 18 patients (45.0%) below 65 years and 22 patients (55.0%) aged 65 or above. According to the Eastern Cooperative Oncology Group Performance Status (ECOG-PS) scale, 16 patients (40.0%) were scored at 0, indicating fully active, able to carry on all pre-disease performance without restriction, whereas 24 patients (60.0%) had a score of 1 or higher, reflecting some limitations in physical activity. Histological analysis revealed that 23 patients (57.5%) had squamous cell carcinoma, 24 (60.0%) had

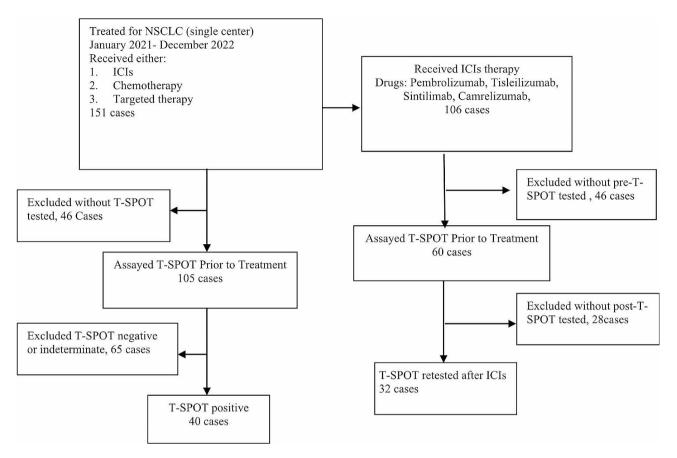


Fig. 1 Flow chart showing cohort selection

adenocarcinoma, and the remaining 3 presented with other histological types. Staging information indicated that 11 patients (27.5%) were at stage III, while a larger group of 29 (72.5%) was at stage IV. Metastasis patterns showed that 8 patients (20.0%) had brain metastasis, 13 (32.5%) had bone metastasis, and 2 (5%) had liver metastasis. Additionally, 19 patients (47.5%) were found to have positive results for driver gene mutations. During the treatment course, targeted therapy was administered to 12 patients (30.0%). It was noted that none of the patients had been diagnosed with active pulmonary tuberculosis prior to treatment initiation.

For the purpose of this study, the patients were divided into two groups based on whether they received ICIs treatment or not: the ICIs treatment group and the non-ICIs treatment group. Comparative analysis between the two groups showed no statistically significant differences in terms of gender, age, stage of cancer, histological type, performance status score, presence of metastasis, or driver gene status (Table 1).

Tuberculosis activity

Among the patients in the ICIs treatment group, a total of 3 cases (11.5%) were diagnosed with active pulmonary tuberculosis during follow-up, occurring at intervals of 2 months, 3 months, and 12 months after ICIs treatment initiation. In contrast, no cases of active pulmonary tuberculosis were observed in the non-ICIs treatment group. However, the difference between the two groups did not reach statistical significance (P=0.186, Table 2).

Impact of ICIs treatment on T-SPOT spot Patient characteristics

A total of 32 NSCLC patients were monitored for T-SPOT tests before and after immunotherapy (Fig. 1). Out of these, 30 (93.8%) were male, while 2 (6.2%) were female. The median age stood at 61 years, with a distribution where 19 patients (59.4%) aged below 65 years and 13 patients (40.6%) were aged 65 years or older. The ECOG-PS results showed an equal distribution, with 16 patients (50.0%) scoring 0 and another 16 patients (50.0%) scoring 1. In terms of histological analysis revealed that 15 patients (46.9%) had squamous cell carcinoma, 15 patients (46.9%) had adenocarcinoma, and 2 patients (6.2%) had adenosquamous carcinoma. As for the disease stage, 11 patients (34.4%) were classified as stage III, while 21 patients (65.6%) were classified as stage IV. The distribution of treatment agents was as follows: pembrolizumab (15.6%), tislelizumab (43.8%), camrelizumab (15.6%), and sintilimab (25.0%). Among the patients,

| Characteristics | All patients (N=40) | ICIs Therapy (<i>n</i> = 26) | Non-ICIs Therapy (<i>n</i> = 14) | P value |
|--------------------------|------------------------|----------------------------------|-----------------------------------|---------|
| Age, years | (| (| | 0.125 |
| <65 | 18(45.0) | 14 | 4 | |
| ≥65 | 22(55.0) | 12 | 10 | |
| Sex | | | | 0.068 |
| Male | 32(80.0) | 23 | 9 | |
| Female | 8(20.0) | 3 | 5 | |
| ECOG-PS | | | | 0.685 |
| 0 | 16(40.0) | 11 | 5 | |
| ≥1 | 24(60.0) | 15 | 9 | |
| Histopathology | | | | 0.179 |
| Adenocarcinoma | 24(60.0) | 13 | 11 | |
| Squamous cell carcinoma | 13(32.5) | 11 | 2 | |
| other | 3(7.5) | 2 | 1 | |
| TNM | | | | 0.170 |
| Stage-III | 11(27.5) | 9 | 2 | |
| Stage-IV | 29(72.5) | 17 | 12 | |
| Metastasis | | | | 0.281 |
| Brain | 8(20.0) | 5 | 3 | |
| Bone | 13(32.5) | 7 | 6 | |
| Liver | 2(5.0) | 0 | 2 | |
| Driver genes Positive | 19(47.5) | 10 | 9 | 0.817 |

| Table 1 | Demographic ar | d clinical d | characteristics of | patients with a | dvanced-stage | lung cancer |
|---------|----------------|--------------|--------------------|-----------------|---------------|-------------|
| | | | | | | |

Table 2 Tuberculosis activity between ICIs and Non-ICIs treatment group

| | ICIs Therapy n = 26 | Non-ICIs Therapy n=14 | P value |
|---------------------|------------------------|--------------------------|---------|
| Active tuberculosis | 3 | 0 | 0.186 |
| Latent infection | 23 | 14 | |

24 (75.0%) received first-line immunotherapy, while 8 (25.0%) received second-line treatment. Brain metastasis was observed in 7 patients (21.9%), bone metastasis in 9 patients (28.1%), and liver metastasis in 2 patients (6.2%). A total of 10 patients (31.2%) tested positive for driver gene mutations, and 3 patients (9.4%) had a history of prior targeted therapy. Importantly, none of the patients were diagnosed with active pulmonary tuberculosis prior to treatment (Table 3).

Changes in T-SPOT and prognostic analysis in NSCLC patients received ICIs treatment

As of May 31, 2023, among the 32 patients, 4 cases showed an increase in spot counts after treatment. Out of these, 2 cases were classified as PD, while no deaths were reported. Additionally, 12 patients had no change in spot counts after treatment, all of whom had a spot count of 0 before treatment. Among this group, 8 patients experienced PD, and 3 patients died, all due to tumor-related causes. Furthermore, 16 patients exhibited a decrease in spot counts after treatment. Among the 8 patients who had spot counts greater than 0 before treatment, 10 experienced PD, and 4 patients died. Notably, one death was attributed to infection with the novel coronavirus, while the remaining deaths were tumor-related causes. The proportions of disease progression in the three groups were 50.0%, 75%, and 62.5%, respectively, with a p-value of 0.837. The proportions of deaths were 0%, 25%, and 25% in the three groups, respectively, with a p-value of 0.527(Table 4).

TB activity and T-SPOT results changes after ICIs treatment During the follow-up period until the cutoff date, 3 cases (9.38%) of active pulmonary tuberculosis were diagnosed among NSCLC patients undergoing immune checkpoint inhibitors (ICIs) treatment. Of these cases, 2 were confirmed positive through sputum tuberculosis smear examination, while 1 case showed detection of mycobacterium tuberculosis via sputum metagenomic next-generation sequencing (mNGS). The onset of active pulmonary tuberculosis was noted at 2 months, 3 months, and 12 months after the initiation of ICIs treatment. Among the 16 patients with a decrease in T-SPOT spot count, 3 cases (18.75%) were diagnosed with active pulmonary tuberculosis. In detail, one case had a decrease in T-SPOT spot count from 16 to 0, another

| Characteristics | | N=32 | (%) |
|------------------|-------------------------|------|------|
| Age, years | | | |
| | <65 | 19 | 59.4 |
| | ≥65 | 13 | 40.6 |
| Sex | | | |
| | Male | 30 | 93.8 |
| | Female | 2 | 6.2 |
| ECOG-PS | | | |
| | 0 | 16 | 50.0 |
| | 1 | 16 | 50.0 |
| Histopathology | | | |
| | Adenocarcinoma | 15 | 46.9 |
| | Squamous cell carcinoma | 15 | 46.9 |
| | Other | 2 | 6.2 |
| TNM | | | |
| | Stage-III | 11 | 34.4 |
| | Stage-IV | 21 | 65.6 |
| ICIs | Pembrolizumab | 5 | 15.6 |
| | Tislelizumab | 14 | 43.8 |
| | Camrelizumab | 5 | 15.6 |
| | Sintilimab | 8 | 25.0 |
| ICIs lines | First line | 24 | 75.0 |
| | Second line | 8 | 25.0 |
| Metastasis | | | |
| | Barin | 7 | 21.9 |
| | Bone | 9 | 28.1 |
| | Liver | 2 | 6.4 |
| Driver genes | Positive | 10 | 31.2 |
| Targeted therapy | | 3 | 9.4 |

| Table 3 | Characteristics of | of 32 NSCLC | patients with | pre- and | post-T-SPOT testing |
|---------|--------------------|-------------|---------------|----------|---------------------|
| | | | | | |

Table 4 Changes in T-SPOT spot counts and prognosis in NSCLC patients undergoing ICIs treatment

| T-SPOT Counts | Increase | No change | Decrease | P value |
|-----------------|----------|-----------|--------------|---------|
| Prognosis, n(%) | n=4 | n=12 | <i>n</i> =16 | |
| PD | 2(50.0) | 8(66.7) | 10(62.5) | 0.527 |
| DCR | 2(50.0) | 4(33.3) | 6(37.5) | |
| Dead | 0(0.0) | 3(25.0) | 4(25.0) | 0.837 |
| Survival | 4(100.0) | 9(75.0) | 12(75.0) | |

case had a decrease from 25 to 4, and the third case had a decrease from 44 to 18. Following diagnosis, all 3 patients temporarily discontinued immune therapy and initiated anti-tuberculosis treatment.

Discussion

A comprehensive meta-analysis has indicated an elevated risk of tuberculosis in cancer patients when compared to the general population. Particularly in the realm of solid tumors, lung cancer stands out with a sixfold increased risk in patients [14]. Both lung cancer and tuberculosis share similar clinical manifestations, and there are cases where they can be diagnosed simultaneously or even be causally linked. Chronic inflammation, genomic changes, and fibrosis associated with tuberculosis are recognized as significant contributors to potential carcinogenic processes. Conversely, the progression of lung cancer and antitumor therapies may also trigger the reactivation of latent tuberculosis infections [15]. Moreover, the presence of untreated inactive tuberculosis lesions in the lungs, coupled with risk factors such as diabetes, smoking, and corticosteroid use, increases the risk for the reactivation of latent tuberculosis [16]. In a study conducted in Japan, among 39 lung cancer patients who subsequently developed tuberculosis (diagnosed more than one month after lung cancer), 37 presented risk factors for tuberculosis or previous tuberculosis infections, with 19 exhibiting two or more such factors [5]. Additionally, the combined use of anticancer drugs with other treatments [17] and the use of ICIs [18] have been linked to the risk of developing tuberculosis.

Previous evidence has suggested a potential association between ICIs treatment and an increased risk of developing pulmonary tuberculosis [19]. In our research, we detected 3 cases (9.38%) of active pulmonary tuberculosis during ICIs treatment. Interestingly, these patients showed a significant decrease in T-SPOT spot counts after ICIs therapy. Yet, the number of spots produced by Phytohemagglutinin (PHA), a non-specific stimulant that induces INF-y release from T cells as a positive control, consistently remained above 100. This indicates a depletion of antigen-specific effector T lymphocytes following ICIs treatment, exposing patients to a dual risk of infection and tumor progression. Notably, one of the 3 cases of active pulmonary tuberculosis in our study succumbed to a severe SARS-CoV-2 infection. International studies have also highlighted an increased prevalence of infectious diseases reliant on T cell-mediated immunity in individuals with reduced T cell counts [10, 19].

Based on the IGRA results before treatment, tuberculosis disease activity can be divided into two categories: newly acquired and reactivation. In a retrospective study, approximately 1.7% (5/297) of lung cancer patients undergoing ICIs treatment developed active pulmonary tuberculosis during the course of therapy. Among these cases, 60% (3/5) were already T-SPOT positive even before initiating ICIs treatment [9]. This observation aligns with similar findings reported in another prospective cohort study, which posits ICIs treatment as a risk factor for the development and exacerbation of tuberculosis [12]. In our study, 3 cases of active tuberculosis were observed, and all had a positive T-SPOT result before treatment initiation, pointing to reactivation of latent tuberculosis infection (LTBI).

The occurrence of active tuberculosis following ICIs treatment may be linked not solely to ICIs-induced T cell depletion but also to the synergistic effects of various factors [20]. For example, the use of corticosteroids during treatment to manage immune-related adverse events may also independently heighten the risk of LTBI reactivation and progression. Furthermore, the prevalence of tuberculosis in a given region can affect the incidence of pulmonary tuberculosis following ICIs treatment. In exploring the discussion of immunological mechanisms, studies have demonstrated that mice lacking the PD-1 receptor are more vulnerable to Mycobacterium tuberculosis [21]. Contrary to the previously held notion that immune dysfunction leads to tuberculosis activation, the disruption of immune tolerance and excessive immune activation can equally play a role in the activation of Mycobacterium tuberculosis [22, 23]. The interaction between PD-1 and PD-L1 creates a state of immune tolerance within the body [24], yet, the use of PD-1 inhibitors in ICIs treatment disrupts this delicate balance. While it stimulates the body's immune response against cancer cells,

thus having an anticancer effect [25]. It may also lead to excessive inflammation and the release of inflammatory cytokines such as TNF- α , IL-18, and IFN- γ , which are implicated in tissue damage and cavity formation. These dynamics can further promote the occurrence and progression of tuberculosis [26].

The T-SPOT spot count in patients after ICIs treatment is determined by both the immune status of T lymphocytes and the burden of Mycobacterium tuberculosis within the body. As ICIs alleviate the inhibitory state of T lymphocytes and enhance their function and quantity, a persistent positive T-SPOT result or a conversion from negative to positive can be critical indicators of tuberculosis activation or reactivation [19]. Our study indicates that when T-SPOT shifts from positive to negative or when there is a decrease in spot count following ICIs treatment, patients may be experiencing antigen-specific activation and depletion of T lymphocytes. Hence, it is imperative to maintain high vigilance for the possibility of tuberculosis activation or reactivation. Routine screening and treatment for latent Mycobacterium tuberculosis infection ought to be conducted before and after ICIs treatment [27].

The implications of ICIs treatment on T-SPOT results remains uncertain, and prudence is advised when interpreting T-SPOT results in patients undergoing ICIs treatment, particularly in cases of T-SPOT conversion to negative or a decrease in spot count. Such changes do not necessarily signify an improvement in tuberculosis infection but may rather serve as a warning sign for tuberculosis activation.

Limitations of this study include its retrospective design and small sample size, which underscore the need for larger-scale, multicenter, or prospective studies to more comprehensively explore this topic. To broaden the research horizon, future studies can extend beyond NSCLC patients receiving ICIs treatment to include a wider range of tumor types, aiming to mitigate potential adverse effects of tuberculosis activation in cancer patients.

Conclusions

In NSCLC patients with a positive T-SPOT response undergoing ICIs treatment, cases of active tuberculosis were observed. Post-ICIs treatment, the changes in T-SPOT spot count demonstrated variability, with a notably higher occurrence of active tuberculosis in patients who experienced a decrease in T-SPOT spot count. This highlights the need to re-evaluate the predictive value of T-SPOT for tuberculosis activity in NSCLC patients treated with ICIs through more extensive research.

Acknowledgements

We gratefully acknowledge the staff, providers and personnel at Zhongshan Hospital(Xiamen), Fudan University who contributed to this study.

Author contributions

YX, QZ, ZC, and HJ contributed to the study conception and design. YX, QZ, and ZC carried out the research, analyzed and interpreted the data, and prepared the manuscript. SY and HC helped collect and organize the date. XX and HJ revised the manuscript. All authors have approved this manuscript for submission.

Funding

This work was supported by grants from Natural Science Foundation of Fujian province (No. 2023J011690), Shanghai science and technology innovation action plan (No. 21140902700), Major medical and health projects of Xiamen (No. 3502Z20204008).

Data availability

The datasets generated during and/or analyzed during the current study are available from the corresponding author on reasonable request.

Declarations

Ethical approval and consent to participate

The study was approved by the ethics committee of Zhongshan Hospital(Xiamen), Fudan University (No. B2022-068). Informed consent from the patients included in this study was not required due to the retrospective nature of the study, and any personal information from the data was deleted beforehand. The study conformed to the provisions of the Declaration of Helsinki (as revised in 2013).

Consent for publication

Not applicable.

Competing interests

The authors declare no competing interests.

Author details

¹Zhongshan Hospital (Xiamen), Fudan University, Xiamen, China ²Xiamen Clinical Research Center for Cancer Therapy, Xiamen, China ³Zhongshan Hospital, Fudan University, Shanghai, China ⁴Fudan Zhangjiang Institute, Shanghai, China

Received: 30 July 2023 / Accepted: 22 March 2024 Published online: 12 April 2024

References

- 1. Bagcchi S. WHO's Global Tuberculosis Report 2022. Lancet Microbe. 2023;4(1):e20.
- Hwang SY, Kim JY, Lee HS, Lee S, Kim D, Kim S, Hyun JH, Shin JI, Lee KH, Han SH et al. Pulmonary tuberculosis and risk of Lung Cancer: a systematic review and Meta-analysis. J Clin Med 2022, 11(3).
- Cabrera-Sanchez J, Cuba V, Vega V, Van der Stuyft P, Otero L. Lung cancer occurrence after an episode of tuberculosis: a systematic review and metaanalysis. Eur Respir Rev 2022, 31(165).
- Ho JC, Leung CC. Management of co-existent tuberculosis and lung cancer. Lung Cancer. 2018;122:83–7.
- Tamura A, Ikeda M, Shimada M, Yamane A. Tuberculosis during lung cancer treatment-A case series. J Infect Chemother. 2022;28(2):339–42.
- Zhou G, Luo Q, Luo S, Teng Z, Ji Z, Yang J, Wang F, Wen S, Ding Z, Li L, et al. Interferon-γ release assays or tuberculin skin test for detection and management of latent tuberculosis infection: a systematic review and meta-analysis. Lancet Infect Dis. 2020;20(12):1457–69.
- Bagchi S, Yuan R, Engleman EG. Immune Checkpoint inhibitors for the treatment of Cancer: clinical impact and mechanisms of response and resistance. Annu Rev Pathol. 2021;16:223–49.
- Tang Q, Chen Y, Li X, Long S, Shi Y, Yu Y, Wu W, Han L, Wang S. The role of PD-1/PD-L1 and application of immune-checkpoint inhibitors in human cancers. Front Immunol. 2022;13:964442.

- Langan EA, Graetz V, Allerheiligen J, Zillikens D, Rupp J, Terheyden P. Immune checkpoint inhibitors and tuberculosis: an old disease in a new context. Lancet Oncol. 2020;21(1):e55–65.
- Kauffman KD, Sakai S, Lora NE, Namasivayam S, Baker PJ, Kamenyeva O, Foreman TW, Nelson CE, Oliveira-de-Souza D, Vinhaes CL et al. PD-1 blockade exacerbates Mycobacterium tuberculosis infection in rhesus macaques. Sci Immunol 2021, 6(55).
- Sakai S, Kauffman KD, Sallin MA, Sharpe AH, Young HA, Ganusov VV, Barber DL. CD4 T cell-derived IFN-gamma plays a minimal role in control of pulmonary Mycobacterium tuberculosis infection and must be actively repressed by PD-1 to prevent Lethal Disease. PLoS Pathog. 2016;12(5):e1005667.
- Fujita K, Elkington P, Redelman-Sidi G, Kanai O, Yamamoto Y, Imakita T, Okamura M, Nakatani K, Mio T. Serial interferon-gamma release assay in lung cancer patients receiving immune checkpoint inhibitors: a prospective cohort study. Cancer Immunol Immunother. 2022;71(11):2757–64.
- Eisenhauer EA, Therasse P, Bogaerts J, Schwartz LH, Sargent D, Ford R, Dancey J, Arbuck S, Gwyther S, Mooney M, et al. New response evaluation criteria in solid tumours: revised RECIST guideline (version 1.1). Eur J Cancer. 2009;45(2):228–47.
- Dobler CC, Cheung K, Nguyen J, Martin A. Risk of tuberculosis in patients with solid cancers and haematological malignancies: a systematic review and meta-analysis. Eur Respir J 2017, 50(2).
- Preda M, Tanase BC, Zob DL, Gheorghe AS, Lungulescu CV, Dumitrescu EA, Stanculeanu DL, Manolescu LSC, Popescu O, Ibraim E et al. The bidirectional relationship between Pulmonary Tuberculosis and Lung Cancer. Int J Environ Res Public Health 2023, 20(2).
- Getahun H, Matteelli A, Abubakar I, Aziz MA, Baddeley A, Barreira D, Den Boon S, Borroto Gutierrez SM, Bruchfeld J, Burhan E, et al. Management of latent Mycobacterium tuberculosis infection: WHO guidelines for low tuberculosis burden countries. Eur Respir J. 2015;46(6):1563–76.
- Kim HR, Hwang SS, Ro YK, Jeon CH, Ha DY, Park SJ, Lee CH, Lee SM, Yoo CG, Kim YW, et al. Solid-organ malignancy as a risk factor for tuberculosis. Respirology. 2008;13(3):413–9.
- Anand K, Sahu G, Burns E, Ensor A, Ensor J, Pingali SR, Subbiah V, Iyer SP. Mycobacterial infections due to PD-1 and PD-L1 checkpoint inhibitors. ESMO Open 2020, 5(4).
- 19. Fujita K, Terashima T, Mio T. Anti-PD1 antibody treatment and the development of Acute Pulmonary Tuberculosis. J Thorac Oncol. 2016;11(12):2238–40.
- Stroh GR, Peikert T, Escalante P. Active and latent tuberculosis infections in patients treated with immune checkpoint inhibitors in a non-endemic tuberculosis area. Cancer Immunol Immunother. 2021;70(11):3105–11.
- 21. Tousif S, Singh Y, Prasad DV, Sharma P, Van Kaer L, Das G. T cells from programmed Death-1 deficient mice respond poorly to Mycobacterium tuberculosis infection. PLoS ONE. 2011;6(5):e19864.
- Elkington PT, Bateman AC, Thomas GJ, Ottensmeier CH. Implications of tuberculosis reactivation after Immune Checkpoint Inhibition. Am J Respir Crit Care Med. 2018;198(11):1451–3.
- Morelli T, Fujita K, Redelman-Sidi G, Elkington PT. Infections due to dysregulated immunity: an emerging complication of cancer immunotherapy. Thorax. 2022;77(3):304–11.
- 24. Boussiotis VA. Molecular and biochemical aspects of the PD-1 checkpoint pathway. N Engl J Med. 2016;375(18):1767–78.
- Tezera LB, Bielecka MK, Ogongo P, Walker NF, Ellis M, Garay-Baquero DJ, Thomas K, Reichmann MT, Johnston DA, Wilkinson KA et al. Anti-PD-1 immunotherapy leads to Tuberculosis reactivation via dysregulation of TNF-alpha. Elife 2020, 9.
- 26. Elkington PT, Friedland JS. Permutations of time and place in tuberculosis. Lancet Infect Dis. 2015;15(11):1357–60.
- Shi J, Li J, Wang Q, Cheng X, Du H, Han R, Li X, Zhao C, Gao G, He Y, et al. The safety and efficacy of immunotherapy with anti-programmed cell death 1 monoclonal antibody for lung cancer complicated with Mycobacterium tuberculosis infection. Transl Lung Cancer Res. 2021;10(10):3929–42.

Publisher's Note

Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.