

Prediction of immune checkpoint inhibitor response in non-small cell lung cancer patients using a microbiome-based biomarker

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INTRODUCTION

- Despite advances in immunotherapy, a significant number of non-small cell lung cancer (NSCLC) patients **do not respond to immune checkpoint inhibitor (ICI) therapies** or **develop immune related adverse events (irAEs)**, leading to treatment discontinuation.
- Recent research indicates that the **microbial signature** may predict **treatment success rates** and thus aid in **optimizing patient selection**.
- BiomeOne[®]** is a **tumor-agnostic CE-IVD** marked medical device that uses stool samples to analyze the **intestinal microbiome (Figure 1)**.
- The aim of our study was to evaluate the **prognostic potential of BiomeOne[®]** in a cohort of NSCLC patients treated with ICI.

RESULTS

- A **total of 42** stage III/IV NSCLC patients (age 48–83 years, mean 64.86 ± 9.33; 59,52% male; 26% in stage III and 74% in stage IV) from **two institutions** were enrolled in this study.
- 100% of the patients received PD1/PDL1 therapy; **50%** of patients had some **immune-related adverse events**.
- The BiomeOne[®] test had an overall **sensitivity of 80,6%** and a **specificity of 54,5%** in predicting the response to ICI therapy in this cohort (see **Table 1**).

Sample classification	Clinical Responder (CR + PR)	Clinical Non-responder (SD + PD)	Total
BiomeOne: Responder	25	5	30
BiomeOne: Non-Responder	6	6	12
Total	31	11	42

Table 1. BiomeOne[®] identified a total of 25/31 clinical responders, and a total of 5/11 non-responders, based on the microbial profile of the baseline stool samples, prior to therapy initiation.

CONCLUSION

The newly developed non-invasive stool test can be used in clinical practice as a predictive test of the ICI response in NSCLC patients. To confirm these results, biomarker analysis in a larger patient cohort is required.

METHODS

- We recruited patients with **unresectable stage III and IV NSCLC** who underwent first line ICI therapy at two institutions.
- Patients who received systemic antibiotic treatment within 30 days prior to treatment initiation were excluded.
- Stool samples were collected prior to ICI therapy using an at-home collection kit.
- After DNA extraction, stool samples underwent **16S rRNA sequencing** and bioinformatic analysis.
- The **microbial profile of each sample** was further analyzed with the proprietary biomarker to **compute the probability of response to ICIs**.
- The patients were classified as **responders (R)** and **non-responders (NR)**. Responders were classified as patients with a radiological and clinical response to the immunotherapy, and non-responders were classified as those with a stable or progressive disease after a 12-week follow-up.



Figure 1. BiomeOne[®] process. A stool collection kit can be ordered by the doctor and delivered directly to the patient. The at-home collection kit is used by the patient to collect the sample and send it directly to the company for sequencing and bioinformatics analysis. The doctor will receive a report with the probability of response to ICI (CE-IVD), probability of developing irAEs (RUO) and exploratory microbiome results.

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