

# Longkankernet implementation ctDNA Analysis

Published: 17-05-2021

Last updated: 04-04-2024

Targeted therapies guided by molecular diagnostics have become a standard treatment of lung cancer. Nowadays, tumour tissue is required to assess the molecular characteristics of the tumour. However, the procedures to obtain tissue are invasive and...

<b>Ethical review</b>	Approved WMO
<b>Status</b>	Recruitment stopped
<b>Health condition type</b>	Respiratory and mediastinal neoplasms malignant and unspecified
<b>Study type</b>	Observational invasive

## Summary

### ID

NL-OMON50777

### Source

ToetsingOnline

### Brief title

LICA

### Condition

- Respiratory and mediastinal neoplasms malignant and unspecified

### Synonym

Lung cancer; non-small cell lung cancer

### Research involving

Human

### Sponsors and support

**Primary sponsor:** Radboud Universitair Medisch Centrum

**Source(s) of monetary or material Support:** 4e geldstroom: commercieel,Astra Zeneca,Hoffmann-La Roche,Novartis

## Intervention

**Keyword:** Circulating tumor DNA, Liquid biopsy, Next-generation sequencing, Non-small cell lung cancer

## Outcome measures

### Primary outcome

The primary aim of this study is to implement the ctDNA analysis to identify actionable targets in lung tumors.

We envision this study will create awareness in regional hospitals with this novel test method.

### Secondary outcome

1. Percentage of patients identified as having an activating and targetable genetic alteration as determined by tissue and liquid biopsies with an intra-individual comparison of the presence of driver mutations detected in tissue and liquid biopsies.
2. To get insight in the time it takes from diagnosis to start of targeted treatment.
3. The percentage of patient with inadequate tissue for analysis where ctDNA may be an alternative.
4. Exploring cost- effectiveness

# Study description

## Background summary

Lung cancer is the second most common cancer and the leading cause of cancer deaths for men and women, In 2019, 9.843 patients were diagnosed with non-small cell lung cancer (NSCLC) in the Netherlands. Despite recent advances made in diagnosis and treatment strategies, NSCLC prognosis remains poor, with a 5-yr overall survival of 15%. In recent years, new methods have been developed for high-throughput molecular analysis of tumours and have provided markers as powerful tools for the development of innovative diagnostic and therapeutic strategies in cancer. Genomic studies have been pioneering in providing information on lung cancer molecular biology, followed by clear evidence that genetic alterations are driving carcinogenesis. Activating somatic mutations including point substitution, small insertion, and in-frame deletion are major oncogenic drivers in lung cancer.

## Study objective

Targeted therapies guided by molecular diagnostics have become a standard treatment of lung cancer. Nowadays, tumour tissue is required to assess the molecular characteristics of the tumour. However, the procedures to obtain tissue are invasive and with a concomitant risk of adverse events. A liquid biopsy, which assess circulating tumor DNA (ctDNA) from peripheral blood, may be an alternative for the majority of NSCLC patients.

Nonetheless, ctDNA analysis is not commonly requested in clinical practice due to limited knowhow, experience, availability or lacking logistics. This study aims to implement ctDNA analysis in daily clinical care by adding the logistics needed for this novel test to the regular clinical practice.

## Study design

The LICA study is a multicenter trial in which ctDNA analysis will be performed using prospectively collected plasma samples.

At the moment of diagnosis, a single blood draw of 25mL will be requested. In the majority of patients, a tissue biopsy will also be performed as part of the regular clinical care. When there is a high suspicion for stage III/IV disease based on imaging, the ctDNA from plasma will be analyzed. The interpretation of the ctDNA analysis will be independent of the biopsy results and will be reported to the treating physician at a later stage.

## Study burden and risks

A sample of blood will be drawn for molecular profiling purposes. This will be

combined with a scheduled blood drawing if possible.

The burden and risks associated with this study are thought to be in balance with the aim to implement this blood based test in the future to replace invasive tissue obtaining procedures.

## Contacts

### Public

Radboud Universitair Medisch Centrum

Geert Grooteplein Zuid 10  
Nijmegen 6525GA  
NL

### Scientific

Radboud Universitair Medisch Centrum

Geert Grooteplein Zuid 10  
Nijmegen 6525GA  
NL

## Trial sites

### Listed location countries

Netherlands

## Eligibility criteria

### Age

Adults (18-64 years)  
Elderly (65 years and older)

### Inclusion criteria

- Patients with confirmed or suspected non-small cell lung cancer
- Signed informed consent
- Adult ( $\geq 18$  years of age) and mentally competent

## Exclusion criteria

- Patients with a benign lesion
- Patients with a metastasis of another primary cancer than lung cancer

## Study design

### Design

**Study type:** Observational invasive

Masking: Open (masking not used)

Control: Uncontrolled

Primary purpose: Diagnostic

### Recruitment

NL

Recruitment status: Recruitment stopped

Start date (anticipated): 07-07-2021

Enrollment: 250

Type: Actual

## Ethics review

Approved WMO

Date: 17-05-2021

Application type: First submission

Review commission: CMO regio Arnhem-Nijmegen (Nijmegen)

Approved WMO

Date: 09-06-2021

Application type: Amendment

Review commission: CMO regio Arnhem-Nijmegen (Nijmegen)

Approved WMO

Date: 28-06-2021

Application type: Amendment

Review commission: CMO regio Arnhem-Nijmegen (Nijmegen)

Approved WMO  
Date: 22-12-2021  
Application type: Amendment  
Review commission: CMO regio Arnhem-Nijmegen (Nijmegen)

## Study registrations

### Followed up by the following (possibly more current) registration

No registrations found.

### Other (possibly less up-to-date) registrations in this register

No registrations found.

### In other registers

Register	ID
CCMO	NL76613.091.21