

# Candida auris screening, surveillance and infection control

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		<input type="checkbox"/> Protocol
<b>Registration date</b> 07/01/2025	<b>Overall study status</b> Ongoing	<input type="checkbox"/> Statistical analysis plan
		<input type="checkbox"/> Results
<b>Last Edited</b> 23/01/2026	<b>Condition category</b> Infections and Infestations	<input type="checkbox"/> Individual participant data
		<input checked="" type="checkbox"/> Record updated in last year

## Plain English summary of protocol

### Background and study aims

Since its discovery in 2009, *Candida auris* cases are rapidly increasing worldwide. In 2017 in Europe, over 600 cases were already reported and in 2021 the United States has reported thousands of cases. In addition to *C. auris*, an increase in other (drug-resistant) *Candida* species is also reported. People most at risk for severe *Candida* infections include those with a weakened immune system, patients undergoing large surgeries and those with indwelling vascular catheters in the intensive care unit (ICU). Because *C. auris* colonizes the human skin and has shown prolonged survival in the environment, it can cause large and persistent outbreaks in hospitals and other healthcare institutions across the globe. In Europe, *C. auris* major outbreaks have been reported in Italy, Spain and the UK, while this species has become endemic in Turkey and Greece. Of concern is that *C. auris* is often multi-drug resistant, leaving the clinician with little to no options for antifungal treatment. For the above reasons, *C. auris* is a major concern for our public health and is categorized as a critical fungal pathogen in the WHO global fungal priority pathogens list. Several cases of *C. auris* colonization have also been found in the Netherlands, but there are currently no active surveillance programs that monitor the epidemiology of (invasive) *Candida* infections. Furthermore, robust screening protocols are lacking, increasing the probability of *C. auris* outbreaks in the Netherlands. Given the rapidly changing epidemiology of *C. auris* and the potential for the emergence of other *Candida* species, it is critical to be prepared for import cases in the Netherlands. This study aims to set up the required infrastructure and protocols to allow early interventions when needed. To combat emerging *Candida* infections, this study aims to build a consortium with stakeholders, evaluate current surveillance systems, and develop optimal screening protocols for *C. auris* in Greece. There will also be work to enhance preparedness with genome sequencing and wastewater surveillance and verify methods through outbreak scenario training. There is overwhelming evidence from medical literature that the epidemiology of yeast is shifting. The results of this project will enable us to implement a search-and-destroy policy concerning *C. auris* and other emerging yeast pathogens, thereby preventing unwanted yeasts from becoming endemic.

### Who can participate?

Patients with a positive yeast culture from blood or other sterile sites (WP2 and 4a); patients in the ICU and patients outside of the ICU that are in contact with *C. auris*-positive patients (WP3); and, patients tested positive for *C. auris* (WP4b)

What does the study involve?

Workpackage 2: Collect 100 yeast isolates from blood in 4 Dutch hospitals and use patient data to assess the ISIS-AR platform for studying invasive yeast in the Netherlands.

Workpackage 3: Screen up to 400 patients in an Athens hospital for *C. auris* at various body sites using different detection methods to determine the best screening protocol.

Workpackage 4A: Use 200 yeast isolates (100 from WP2 and 100 from additional sites) to establish a whole genome sequencing platform for rapid identification and genotyping of yeast, including *C. auris*.

Workpackage 4B: Evaluate the suitability of wastewater sampling for *C. auris* surveillance using isolates from routine MDRO screening in 4 Dutch hospitals.

What are the possible benefits and risks of participating?

In the retrospective part of the study, there are no benefits for the participants. In the prospective portion of the study conducted in Greece, patients might test positive for *Candida auris*, which may not have been identified otherwise. This could result in a more adequate treatment. Additionally, identifying positive cases enables the implementation of infection control measures, helping to prevent the further spread of *C. auris* to other patients.

There are no risks related to participation in the study.

Where is the study run from?

Netherlands National Institute for Public Health and the Environment, RIVM

When is the study starting and how long is it expected to run for?

May 2024 to May 2028

Who is funding the study?

1. ZonMw, Netherlands Organisation for Health Research and Development
2. Netherlands National Institute for Public Health and the Environment, RIVM

Who is the main contact?

Dr Eelco Meijer, [Eelco.meijer@cwz.nl](mailto:Eelco.meijer@cwz.nl)

## Contact information

### Type(s)

Public, Scientific, Principal investigator

### Contact name

Dr Eelco Meijer

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## Additional identifiers

### Clinical Trials Information System (CTIS)

Nil known

### ClinicalTrials.gov (NCT)

Nil known

### Protocol serial number

10150022310025

## Study information

### Scientific Title

Candida AUris screening, surveillance and infectTION control

### Acronym

CAUTION

### Study objectives

Infections with yeasts such as Candida auris have been rising worldwide, including in Europe. Dutch hospitals had multiple C. auris colonized patients recently, all imported. Infections with C. auris mainly affect immunocompromised patients and patients who underwent complex surgeries. Since C. auris can survive well on surfaces, this yeast can spread in healthcare facilities and cause outbreaks. C. auris is often resistant to many antifungal medications hampering the treatment of patients. These factors have contributed to listing C. auris on the WHO fungal priority list, in addition to C. parapsilosis, C. tropicalis and C. albicans. Since screening and surveillance are not well developed, the overview of the presence and spread of C. auris is limited. This project will develop accurate screening and surveillance of yeasts to be prepared for the emergence of C. auris and to be able to adequately respond if C. auris and other yeasts spread and cause outbreaks in the Netherlands.

### Ethics approval required

Ethics approval required

### Ethics approval(s)

1. approved 10/04/2024, Scientific Council of Attikon University Hospital (Rimini 1, Haidari, 12462, Greece; +30 210-583-1692; greps@attikonhospital.gr), ref: 226/22-3-2024

2. approved 22/09/2024, METC Oost-Nederland (Philips van Leydenlaan 25, Nijmegen, 6525EX, Netherlands; +31 024 3613154; METCoost-en-CMO@radboudumc.nl), ref: 2024-17538

### Study design

Multicenter cross-sectional observational study

### Primary study design

Observational

### **Study type(s)**

Diagnostic, Screening

### **Health condition(s) or problem(s) studied**

Candidiasis

### **Interventions**

Workpackage 2: 100 yeast isolates from blood, found during routine diagnostics in 4 Dutch hospitals, will be collected. The isolates, together with some patient information (age, gender) will be used to investigate whether the existing ISIS-AR platform provides sufficient data to study invasive yeast in the Netherlands.

Workpackage 3: To determine the best *C. auris* screening protocol including sampling method and body site, a maximum of 400 patients from a hospital in Athens, Greece, will be screened for *C. auris* at different body sites. The swabs will be analyzed using different detection methods.

Workpackage 4A: The 100 yeast isolates from WP2, together with an additional 100 yeast isolates from blood and other sterile sites collected at 2 more Dutch hospitals, will be included. These isolates will be used to set up a whole genome sequencing platform for rapid identification and genotyping of yeast isolates, including *C. auris*.

Workpackage 4B: The suitability of wastewater sampling for the surveillance of *C. auris* will be evaluated. *C. auris* isolates found during routine MDRO screening at the 4 participating Dutch hospitals will be used in the study to confirm if *C. auris* is present in the hospitals' wastewater.

### **Intervention Type**

Not Specified

### **Primary outcome(s)**

WP2: Determination and susceptibility testing is performed on the yeast isolates found in blood, using the hospitals' own protocols. The isolates will also be sent to the national reference center to verify the results using EUCAST.

WP3: The best body locations for screening of *C. auris* will be tested by taking swabs from different body sites. To determine the best method for the detection of *C. auris*, different methods for culturing and molecular detection (PCR and LAMP) are compared using the swabs.

WP4a: Determination and susceptibility testing will be done in the same manner as for WP2. The isolates collected for WP2 and WP4 will also be sequenced by the national reference center using Illumina sequencing.

WP4B: Wastewater samples will be collected once a month in each center, serving as background data, and partially as a negative control. When proven *C. auris* colonized patients are admitted, wastewater sampling can be intensified.

### **Key secondary outcome(s)**

WP4B: All *C. auris* isolates found during MDRO screening at the participating hospitals will be sent to the national reference center for determination, susceptibility testing using EUCAST, and whole genome sequencing using Illumina sequencing.

### **Completion date**

01/05/2028

### **Eligibility**

**Key inclusion criteria**

WP2 & WP4a: Patients with a positive yeast culture from blood or other sterile sites.

Added 23/01/2026: Inclusion was performed retrospectively and was completed in November 2025.

WP3: Patients in the ICU and patients outside of the ICU who were in contact with C. auris-positive patients.

Added 23/01/2026: Inclusion was completed in January 2026.

WP4b: Patients tested positive for C. auris.

Added 23/01/2026: Inclusion is still ongoing.

**Participant type(s)**

Patient

**Healthy volunteers allowed**

No

**Age group**

All

**Sex**

All

**Total final enrolment**

0

**Key exclusion criteria**

Patients who object to participation.

**Date of first enrolment**

01/11/2024

**Date of final enrolment**

01/05/2028

**Locations****Countries of recruitment**

Greece

Netherlands

**Study participating centre**

Amsterdam UMC

Meibergdreef 9

Amsterdam

Netherlands

1105 AZ

**Study participating centre**

**Erasmus MC**

Dr. Molewaterplein 40

Rotterdam

Netherlands

3015GD

**Study participating centre**

**Radboudumc**

Geert Grooteplein Zuid 10

Nijmegen

Netherlands

6525GA

**Study participating centre**

**University Medical Center Groningen**

Hanzeplein 1

Groningen

Netherlands

9713GZ

**Study participating centre**

**St. Antonius Ziekenhuis**

Koekoekslaan 1

Nieuwegein

Netherlands

3435CM

**Study participating centre**

**Laboratorium Microbiologie Twente Achterhoek**

Boerhaavelaan 59

Hengelo

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**Study participating centre**

**University General Hospital 'Attikon'**

Rimini 1

Haidari, Athens  
Greece  
12462

## Sponsor information

### Organisation

National Institute for Public Health and the Environment

### ROR

<https://ror.org/01cesdt21>

## Funder(s)

### Funder type

Not defined

### Funder Name

ZonMw

### Alternative Name(s)

Netherlands Organisation for Health Research and Development

### Funding Body Type

Private sector organisation

### Funding Body Subtype

Other non-profit organizations

### Location

Netherlands

### Funder Name

Rijksinstituut voor Volksgezondheid en Milieu

### Alternative Name(s)

Netherlands National Institute for Public Health and the Environment, RIVM

### Funding Body Type

Government organisation

### Funding Body Subtype

National government

**Location**

Netherlands

## **Results and Publications**

**Individual participant data (IPD) sharing plan**

The datasets generated during and/or analyzed during the current study will be available upon request.

Findable: Data will be available upon request.

Accessible: There will be restrictions to access the data collection. The researchers involved in the project have first been given the opportunity to complete the analyzes and publish the results. After publication, the data in question will be made publicly available. If, after completion of the analyzes and publications, data from the bio-databases and other data have not been made public, a committee in office at that time will decide on further accessibility. Matters such as privacy of data, relevance, intent, commercial approach, conflicts of interest, etc. will be examined.

Interoperable: The data will be stored in such way (adapted to generic open standards) that other researcher can read the data collection.

Re-usable: All project participants have to give their permission for reuse of the data, and the data will be pseudonymised.

**IPD sharing plan summary**

Available on request