

Facility Instructions for *C. auris* Surveillance and Reporting

Review reports from the past 6 months and for the next 3 months for any of the following:

- *Candida auris*
- *Candida haemulonii*
- *Candida duobushaemulonii*
- *Rhodotorula glutinis* (characteristic red color not present)
- *Candida sake*
- *Candida intermedia*
- *Saccharomyces kluyveri*
- *Candida catenulate*
- *Candida famata*
- *Candida guilliermondii*
- *Candida lusitaniae*
- *Candida parapsilosis*

If any of the above are found:

- Determine lab identification method
- Refer to [CDC table](#) to determine if method is valid for identifying *C. auris*

Identification Method	Organism <i>C. auris</i> can be misidentified as
Vitek 2 YST*	<i>Candida haemulonii</i> <i>Candida duobushaemulonii</i>
API 20C	<i>Rhodotorula glutinis</i> (characteristic red color not present) <i>Candida sake</i>
API ID 32C	<i>Candida intermedia</i> <i>Candida sake</i> <i>Saccharomyces kluyveri</i>
BD Phoenix yeast identification system	<i>Candida haemulonii</i> <i>Candida catenulata</i>
MicroScan	<i>Candida famata</i> <i>Candida guilliermondii</i> ** <i>Candida lusitaniae</i> ** <i>Candida parapsilosis</i> **
RapID Yeast Plus	<i>Candida parapsilosis</i> **

- Reference to *C. auris* identification methods can be found on the CDC website: <https://www.cdc.gov/fungal/candida-auris/identification.html>.
- If possible misidentification, the facility might be asked to send an isolate to the DSHS Austin lab to ensure correct identification. Please reach out to your local health department epidemiology contact and they will help you coordinate the submission.