Procedure for identification of yeasts using MALDI-ToF

1.0 Purpose

This document outlines the procedure for identification of yeast isolates using the MALDI-ToF MS Bruker Biotyper platform and a custom extraction protocol optimized to identify isolates that are represented in the Bruker database and the MicrobeNet database.

2.0 Definitions

Term	Definition
MALDI-ToF	Matrix-assisted laser desorption/ionization – time of flight
SAB Media	Sabouraud's Dextrose agar
Chromagar Media	CHROMagar Candida Chromogenic agar
BTS	Bacterial Test Standard

3.0 Equipment: Properly maintained and calibrated per institutions protocols

- 3.1. Biological Safety Cabinet
- 3.2. MALDI-ToF MS Bruker Biotyper

4.0 Reagents/Supplies/Media

- 4.1. 100% Ethanol-Sterile Solution: example, SIGMA (Catalog # 459836), or equivalent. Always use Ethanol labeled "molecular biology reagent."
- 4.2. Sterile WFI Quality Cell Culture Grade Water: example, Fisher Scientific (Product # L4321849), or equivalent
- 4.3. Sabouraud's Dextrose Agar (Emmons): Plates and slants; example, Fisher Scientific (Product #L21827), or equivalent
- 4.4. CHROMagar Candida Chromogenic agar: available from multiple vendors as plates or powder
- 4.5. Reusable polished steel MALDI target plate with 96 sample positions (Bruker Product # 8280800)
- 4.6. Bruker matrix HCCA (HCCA = α -Cyano-4-hydroxycinnamic acid): (Product # 255344) --Prepare by adding 250 µl of Bruker Standard Solvent, vortex until liquid is clear.
- 4.7. 70% Formic Acid: example, SIGMA (Catalogue # 27001), diluted using sterile WFI Quality cell culture grade water
- 4.8. Bacterial test standard (BTS): Bruker (Product # 255343) -- Prepare BTS according to manufacturer's instructions and store at ≤ -18°C until ready for use.

- 4.9. Bruker Standard Solvent (Sigma Aldrich #900666)
- 4.10. 1µl Culture loops
- 4.11. 1.5 mL microcentrifuge tubes

5.0 Safety: All institutional safety procedures must be followed in the performance of this standard operating procedure.

- 5.1. Standard personal protective equipment should follow institutional guidelines but should consist of at least gowns, gloves, and safety glasses, which should be worn at all times.
- 5.2. All of the set-up procedures should be performed in a biological safety cabinet.

5.3. Chemical Safety

- 5.3.1. Formic Acid: Flammable, Toxic, and Corrosive
- 5.3.2. α-Cyano-4-hydroxycinnamic acid (HCCA): Skin irritant, Serious eye irritant.

6.0 Sample Preparation Information/Processing

6.1. Growth

- 6.1.1. Overnight growth of <u>a pure yeast culture</u> should be used for routine microorganism identification;
- 6.1.2. Slow-growing yeast (e.g., *Candida haemulonii*) may need to grow for several days before testing.

6.2. Storage

- 6.2.1. Store plates at room temperature for 1-5 days is acceptable;
- 6.2.2. Culture medium (Sabouraud dextrose agar or CHROMagar) and growth temperature (25°C vs 37°C) have little to no effect on results.

NOTE: <u>Do not</u> use organisms that have been stored at 4°C or lower as this has a negative impact on quality of spectra and reproducibility.

7.0 Quality Control

7.1. For each run, include pure culture of *Candida auris* and *Candida parapsilosis*, in duplicate, as positive controls.

8.0 Procedure

NOTE: Prior to beginning the procedure, note on a sample key spreadsheet (Appendix A) the location of where each specimen will be placed on the 96-spot reusable stainless steel target plate.

8.1. Extraction Procedure

NOTE: <u>Perform the following steps in a biological safety cabinet</u> to ensure safety and sterility.

- 8.1.1. Perform extraction procedure in duplicate.
- 8.1.2. Label two 1.5 mL microcentrifuge tubes for each sample, including QC strains from Step 6.1.
- 8.1.3. Add 50 μ L of sterile WFI quality water to each tube.
- 8.1.4. Using a sterile, clear 1 μl inoculating loop, transfer 1 loopful of microorganism to both tubes containing 50 μl of sterile water. If the colonies are small, transfer more than one colony, choosing isolated colonies, when possible.
- 8.1.5. Secure the lid of the tube.
- 8.1.6. Add **50** μ L of 100% Ethanol to each tube.
- 8.1.7. Using a 100 μl pipette set to 50 μl, pipette the liquid up and down 10 times to homogenize.
- 8.1.8. From each tube, add one microliter of the resulting suspension to a 96-spot reusable stainless steel target plate (see Fig. 1).
- 8.1.9. Let the plate air dry for two minutes.
- 8.1.10. Overlay dried spot with 1 µL of 70% Formic Acid.
- 8.1.11. Air dry for two minutes.
- 8.1.12. Add 1 μL of the BTS suspension to two open spots on the 96-spot reusable stainless steel target plate (<u>do not</u> add formic acid to BTS! Simply allow to air dry and proceed to step 8.1.13). Perform this step once a week. For preparation of BTS, see step 4.8.
- 8.1.13. As soon as spots are dry, immediately overlay with 1 μL of HCCA matrix.
 NOTE: Do not delay the addition of HCCA matrix as it will decrease the quality of your run.
- 8.1.14. Air dry for two minutes.
- 8.1.15. As soon as spots are dry, they are ready to be subject to MALDI-TOF analysis.
 NOTE: target <u>must be completely dry</u> before it is inserted into MALDI BioTyper.
- 8.1.16. Once the target plate is dry, the plate <u>must be run within 24 hours</u>.
- 8.1.17. Be sure that the location of each specimen on the 96-spot reusable stainless steel target plate is recorded on a sample key spreadsheet. Verify the spreadsheet numbers to make sure they match the original sample numbers.
- 8.1.18. Samples are now ready for the MALDI-ToF MS assay on the Bruker Biotyper platform.
- 8.1.19. Resulting spectra must be interpreted using Bruker database.

9.0 Performance Specifications

9.1. Bruker allows for any score above 2.0 to be acceptable for a species identification. A laboratory may perform an in-house validation to determine a lower score which may be acceptable for species identification.

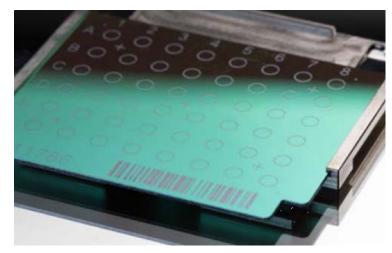


Figure 1. Example of stainless-steel target plate

10.0 Interpretation of Results

10.1. Review control strains.

- 10.1.1. At least one of the duplicates for each control strain must generate a score of ≥ 2 with a correct identification (no conflicting results between the duplicates).
- 10.1.2. If the control strains generate scores < 2 or incorrect identification, the results cannot be interpreted, and the extraction/submission process must be repeated.
- 10.2. *Candida* species will be identified if database match scores are ≥ 2 for one or more of the duplicates.
- 10.3. If no species identification is made, repeat extraction/submission process using a new subculture.
- 10.4. If no identification is successful after second submission, the isolate must be identified by another methodology.

11.0 Procedure for submission of Main Spectra (MSP) to MicrobeNet (Optional)

- 11.1. Go to the MicrobeNet website <u>www.microbenet.cdc.gov.</u>
- 11.2. Request a MicrobeNet account.
- 11.3. Log into account.
- 11.4. Convert Main Spectra to XML file on your Bruker software.
- 11.5. Drag and drop the XML file onto the MicrobeNet website.
- 11.6. The top 10 matches will populate.

12.0 References

12.1. Fraser, M., et al. (2016). "Rapid identification of 6328 isolates of pathogenic yeasts using MALDI-ToF MS and a simplified, rapid extraction procedure that is compatible with the Bruker Biotyper platform and database." Medical Mycology 54(1): 80-88.

Disclaimer

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