



Certificate of Analysis: Lyophilized Microorganism Specification and Performance Upon Release

<b>SPECIFICATIONS:</b> <b>Product Name:</b> Saccharomyces cerevisiae <b>Catalog Number:</b> 0699 <b>Lot Number:</b> 699-252** <b>Reference Number:</b> ATCC® 9763™* <b>Passage from Reference:</b> 3 <b>Expiration Date:</b> 2025/08/31	<b>RELEASE INFORMATION:</b> <b>Quality Control Technologist:</b> Kavitha Gobalan <b>Release Date:</b> 2023/10/06
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<b>Performance</b>	
<b>Macroscopic Features:</b> Medium to large, circular, dull, white to cream colonies.  <b>Microscopic Features:</b> Gram positive, yeast cells, oval to spherical, spores are gram negative when present.	<b>Medium:</b> SAB DEX EMMONS  <b>Method:</b> Gram Stain (1)
<b>ID System:</b> MALDI-TOF (1)	
See attached ID System results document.	

Amanda Kuperus  
Director of Quality Control  
AUTHORIZED SIGNATURE

\*\*Disclaimer: The last digit(s) of the lot number appearing on the product label and packing slip are merely a packaging event number. The lot number displayed on this certificate is the actual base lot number.

**Refer to the enclosed product insert for instructions, intended use and hazard/safety information.**

Individual products are traceable to a recognized culture collection.

(1) These tests are accredited to ISO/IEC 17025.



TESTING CERT #2655.01



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REFERENCE MATERIAL PRODUCER  
CERT #2655.02

# Bruker Daltonik MALDI Biotyper Classification Results



## Meaning of Score Values

Range	Interpretation	Symbols	Color
2.00 – 3.00	High-confidence identification	(+++)	Green
1.70 – 1.99	Low-confidence identification	(+)	Yellow
0.00 – 1.69	No Organism Identification Possible	(-)	Red

## Meaning of Consistency Categories (A - C)

Category	Interpretation
(A)	<b>High consistency:</b> The best match is a high-confidence identification. The second-best match is (1) a highconfidence identification in which the species is identical to the best match, (2) a low-confidence identification in which the species or genus is identical to the best match, or (3) a non-identification.
(B)	<b>Low consistency:</b> The requirements for high consistency are not met. The best match is a high- or low-confidence identification. The second-best match is (1) a high- or low-confidence identification in which genus is identical to the best match or (2) a non-identification.
(C)	<b>No consistency:</b> The requirements for high or low consistency are not met.

Run Creation Date/Time: 2023-10-05T11:36:52.251 KG

Applied MSP Library(ies): BDAL, Mycobacteria Library (bead method), Filamentous Fungi Library

Sample Name	Sample ID	Organism (best match)	Score Value
F12 (+++) (A)	699-252	Saccharomyces cerevisiae	2.06

Comments:

n/a