

## **CANDIDA ALBICANS DRUG RESISTANCE PANEL**

ATCC *Candida albicans* Drug Resistance (CaDR) Panel (ATCC<sup>®</sup> MP-8<sup>™</sup>) represents strains resistant to one or more antifungal drugs. It also includes two sensitive strains that have been typed and sequenced to serve as controls. Strain difference in genetic background is denoted by variance in the yeast DNA barcoding region. This unique set of *Candida albicans* strains is useful for assav development, novel drug testing, and other applications.

| ATCC <sup>®</sup> No. | Strain<br>Designation | D1D2<br>Variance**         | Isolation<br>Source                         | Geographical<br>Region | Anidula-<br>fungin | Mica-<br>fungin | Caspo-<br>fungin | 5-flucy-<br>tosine | Voricon-<br>azole | ltracon-<br>azole | Flucon-<br>azole |
|-----------------------|-----------------------|----------------------------|---------------------------------------------|------------------------|--------------------|-----------------|------------------|--------------------|-------------------|-------------------|------------------|
| <u>64124</u> ™        | Darlington            | T455C                      | Human<br>mouth swab                         | Unknown                | R                  | R               | R                | I                  | R                 | R                 | R                |
| <u>10231</u> ™        | 3147                  | T455C,<br>č601A            | Human<br>bronchomy-<br>cosis                | Unknown                | R                  | S               | S                | S                  | R                 | R                 | R                |
| <u>76485</u> ™        |                       | T455C                      | Human<br>infant<br>cerebrospi-<br>nal fluid | Dallas, Texas<br>(USA) | R                  | S               | R                | S                  | S                 | S                 | S                |
| <u>28121</u> ™        | 304                   | C263T,<br>T455C,<br>~601TT | Human<br>feces                              | California<br>(USA)    | S                  | S               | S                | S                  | R                 | R                 | R                |
| <u>90819</u> ™        | CATW 4/19             | A570G,<br>~601T            | Human                                       | Milwaukee,<br>WI (USA) | S                  | S               | S                | S                  | R                 | R                 | R                |
| <u>MYA-1023</u> ™     | GT 157                | T455C,<br>A570G,<br>~601A  | Human                                       | West Virginia<br>(USA) | S                  | S               | S                | S                  | R                 | R                 | R                |
| <u>MYA-427</u> ™      | A39                   | None                       | Human<br>blood                              | Belgium                | S                  | S               | S                | S                  | R                 | R                 | R                |
| <u>MYA-574</u> ™      | Gu5                   | T455C                      | Human AIDS<br>patient                       | Germany                | S                  | S               | S                | S                  | R                 | R                 | R                |
| <u>38289</u> ™        | Tu 62823              | T455C                      | Human<br>female<br>urethra                  | Unknown                | S                  | S               | S                | S                  | R                 | R                 | S                |
| <u>11651</u> ™        | 171D                  | T455C                      | Human lung<br>pus                           | Virginia (USA)         | S                  | S               | S                | S                  | S                 | R                 | S                |
| <u>96901</u> ™        | 321182                | T455C                      | Human<br>HIV-positive<br>patient            | Omaha, NE<br>(USA)     | S                  | S               | S                | S                  | S                 | S                 | R                |
| <u>90029</u> ™        | NCCLS 67              | T455C                      | Human                                       | lowa (USA)             | S                  | S               | S                | R                  | S                 | S                 | S                |

| ATCC <sup>®</sup> No. | Strain<br>Designation | D1D2<br>Variance** | Isolation<br>Source                           | Geographical<br>Region | Anidula-<br>fungin | Mica-<br>fungin | Caspo-<br>fungin | 5-flucy-<br>tosine | Voricon-<br>azole | ltracon-<br>azole | Flucon-<br>azole |
|-----------------------|-----------------------|--------------------|-----------------------------------------------|------------------------|--------------------|-----------------|------------------|--------------------|-------------------|-------------------|------------------|
| <u>MYA-2876</u> ™     | SC5314*               | T455C              | Human<br>clinical<br>isolate                  | Unknown                | S                  | S               | S                | S                  | S                 | S                 | S                |
| <u>18804</u> ™        | CBS 562               | Type strain        | Human skin<br>- erosio<br>interdigi-<br>talis | Uruguay                | S                  | S               | S                | S                  | S                 | S                 | S                |

\*ATCC MYA-2876''', SC5314, is a widely studied, genome-sequenced, wild type strain that has been included as a control for both gene sequence and drug sensitivity.

\*\*0102 variance refers to the nucleotide difference in the 0102 region of LSU rONA as compared to the Type Strain of *Candida albicans* (ATCC® <u>18804</u>"). The nucleotide position is referenced using GenBank NW\_139715 (C. *albicans* SC5314 28S rRNA gene) annotation. A nucleotide substitution at position 455 from T (in the Type Strain) to C is expressed as T455C. The v symbol denotes an insertion at a given position, e.g., 601 TT denotes an insertion of two T's at position 601.

The data above was generated by ATCC through testing of the earliest seed stock available based on protocols described in: Clinical and Laboratory Standards Institute (CLSI) published standard, Reference Method for Broth Dilution Antifungal Susceptibility Testing of Yeasts; Approved Standard-Second Edition /M27-A2). Analysis was performed using YeastOne® plates Y09 and YeastOne® broth Y3462 from TREK Diagnostic Systems (part of Thermo Fisher Scientific). The interpretation of resistance (R), intermediate /I}, and susceptibility[S] was made according to guidelines described in the vendor's product instruction and relevant CLSI documents. ATCC provides these data in good faith, but makes no warranty, express or implied, nor assumes any legal liability or responsibility for any purpose for which the data are used. The ATCC trademark and trade name, any and all ATCC catalog numbers, and any other trade-marks listed are trademarks of the American Type Culture Collection unless indicated otherwise. YeastOne® is a registered trademark of TREK Diagnostic Systems (part of Thermo Fisher Scientific). ATCC products are intended for laboratory research only. They are not intended for use in humans, animals or diagnostics.



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