

Deletion of *AIF1* but not of *YCA1/MCA1* protects *Saccharomyces cerevisiae* and *Candida albicans* cells from caspofungin-induced programmed cell death

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ABSTRACT Caspofungin was the first member of a new class of antifungals called echinocandins to be approved by a drug regulatory authority. Like the other echinocandins, caspofungin blocks the synthesis of $\beta(1,3)$ -D-glucan of the fungal cell wall by inhibiting the enzyme, $\beta(1,3)$ -D-glucan synthase. Loss of $\beta(1,3)$ -D-glucan leads to osmotic instability and cell death. However, the precise mechanism of cell death associated with the cytotoxicity of caspofungin was unclear. We now provide evidence that *Saccharomyces cerevisiae* cells cultured in media containing caspofungin manifest the classical hallmarks of programmed cell death (PCD) in yeast, including the generation of reactive oxygen species (ROS), the fragmentation of mitochondria, and the production of DNA strand breaks. Our data also suggests that deleting *AIF1* but not *YCA1/MCA1* protects *S. cerevisiae* and *Candida albicans* from caspofungin-induced cell death. This is not only the first time that *AIF1* has been specifically tied to cell death in *Candida* but also the first time that caspofungin resistance has been linked to the cell death machinery in yeast.

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INTRODUCTION

Defined as any cell death that results from the activation of a genetic program, programmed cell death (PCD) has been classified by the Nomenclature Committee on Cell Death (NCCD) into twelve different functional categories based on measurable biochemical features [1]. In recent years, numerous laboratories have shown that a wide range of antifungal drugs leads to PCD in yeast that is reminiscent of apoptosis in mammalian cells [2]. This cell death is accompanied by several distinctive phenotypical markers, including the generation of reactive oxygen species (ROS), the fragmentation of mitochondria, and the dissipation of the mitochondrial membrane potential [3-5]. It also involves intrinsic pathways of cell death defined by several genes, including *AIF1*, *BIR1*, *BX11*, *NMA111*, and *YCA1*, which have homologs in the metazoa [6-11].

Caspofungin was the first member of the class of antifungals called the echinocandins to be approved by a drug

regulatory authority [12, 13]. Like the other echinocandins, caspofungin blocks the synthesis of $\beta(1,3)$ -D-glucan of the fungal cell wall by inhibiting the enzyme, $\beta(1,3)$ -D-glucan synthase [14]. However, the precise mechanism of cell death associated with the cytotoxicity of caspofungin was unclear.

In this report, we provide evidence that *Saccharomyces cerevisiae* cells cultured in media containing caspofungin manifest the classical hallmarks of PCD in yeast, including the generation of reactive oxygen species (ROS), the fragmentation of mitochondria, and the activation of caspase-like activity. Moreover, our data also suggests that this cell death scenario requires the pro-apoptotic gene, *AIF1*, but not the metacaspase, *YCA1*, in both *S. cerevisiae* and *Candida albicans*. Our findings reveal an additional dimension of caspofungin's mechanism of action as a general antifungal agent that points to a possible novel pathway to drug resistance.

RESULTS AND DISCUSSION

To determine if caspofungin triggers PCD in budding yeast, we cultured W303-1A cells in YPD containing 0.02µg/ml caspofungin for three hours, a drug concentration below the MIC of 0.04µg/ml for this strain (data not shown). With

this protocol, propidium iodide staining (50µg/ml) revealed that just over half of the cells (58±10%; p<0.005) were viable as compared to cells grown in the absence of drug (99±0.1%). The cells were also treated with 2.5µM CellROX Green (Molecular Probes, Inc.) and imaged with a Zeiss

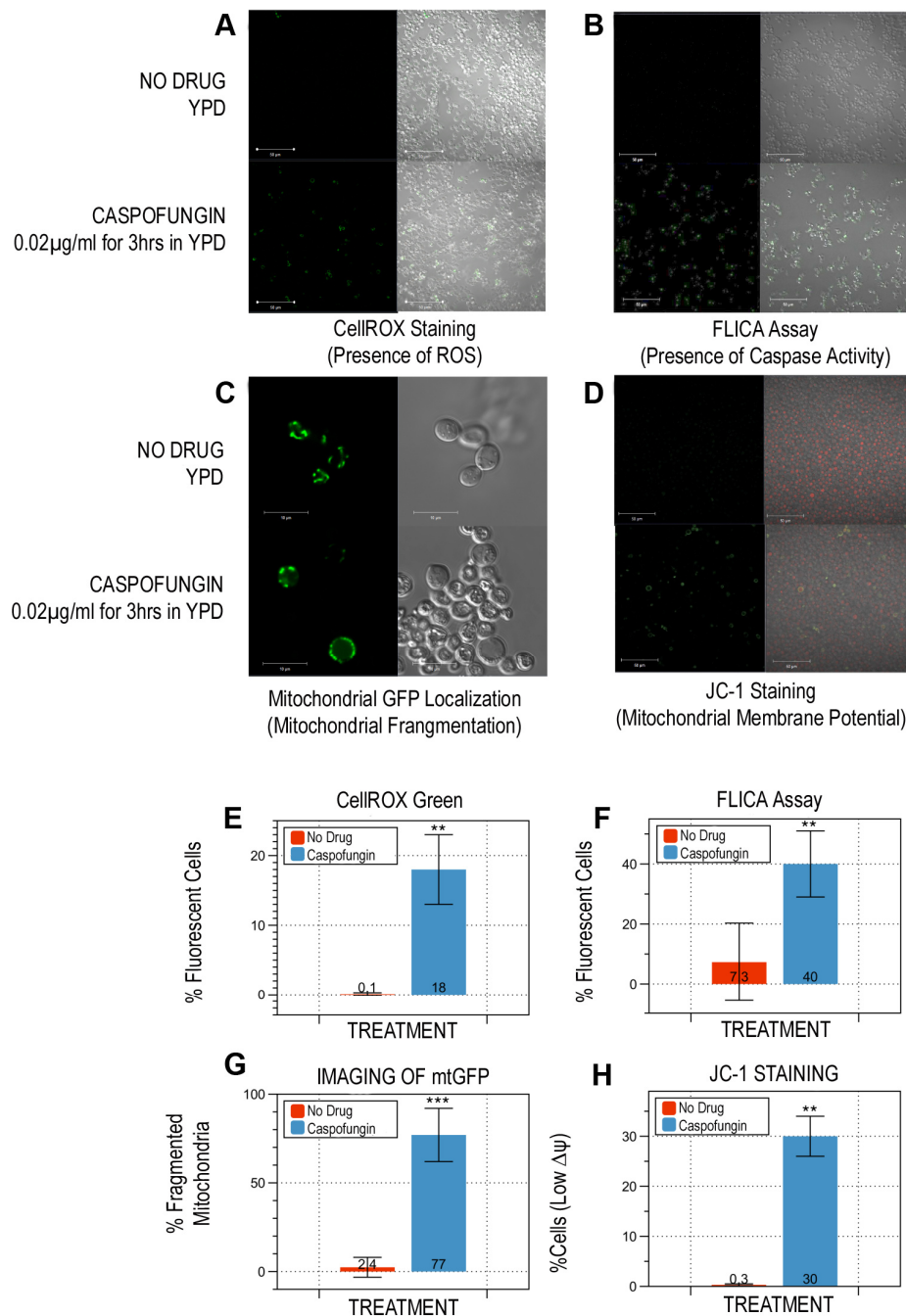


FIGURE 1: *Saccharomyces cerevisiae* cells cultured in caspofungin manifest numerous phenotypic markers associated with yeast programmed cell death. Exponentially-growing wildtype cells from the W303-1A strain background were cultured in YPD containing 0.02µg/ml caspofungin for three hours. The cells were stained with 2.5µM CellROX Green to detect ROS (A, E); with a FLICA kit to detect activated caspase activity (B, F); and with the mitochondrial vital dye, JC-1, to assess their mitochondrial membrane potential (D, H). To visualize the structure of their mitochondria, the cells were transformed with a p416 GPD-mtGFP plasmid constitutively expressing a mitochondria-localized GFP marker and their mitochondria were visualized after they had been treated with caspofungin as described above (C, G). Representative images are shown (A-D). Statistical significance for the graphs (E-H) was determined with the unpaired Student’s t-test (**: p<0.05; ***: p<0.005), and a minimum of 500 cells was counted for each assay.

LSM700 confocal microscope. We discovered that including caspofungin in the culture media generates significantly more fluorescent cells ($18\pm 5\%$; $p < 0.05$; Figure 1A, E) as compared to controls ($0.1\pm 0.2\%$) suggesting that the drug induces the production of ROS, a characteristic hallmark of apoptotic cell death in yeast [3, 15].

Next, we used a FLICA apoptosis detection kit (ImmunoChemistry Technologies, LLC) according to the manufacturer's specifications to determine if the caspofungin treatment described above also induced caspase activity in W303-1A cells. Numerous studies have associated caspase activation with apoptotic cell death, not only in yeast, but also in mammalian cells [16-18]. We discovered that drug treatment leads to significantly more cells with activated caspase-like activity ($40\pm 11\%$; $p < 0.05$; Figure 1B, F) as compared to no-drug controls ($7.3\pm 13\%$).

Third, we imaged the same yeast cells, this time transformed with the p416 GPD-mtGFP plasmid constitutively expressing a mitochondria-localized GFP marker to visualize the structure of their mitochondria [19], after they had been treated with caspofungin as described above. Several reports have shown that apoptosis in *Saccharomyces cerevisiae* is accompanied by the fragmentation of mitochondria [4, 20]. Our experiments revealed that the majority of mitochondria of the caspofungin-treated cells had lost

their tubular architecture after three hours ($77\pm 15\%$; $p < 0.005$; Figure 1C, G). In contrast, only a few cells cultured in media without the drug ($2.4\pm 5.6\%$) displayed the punctate mitochondrial-staining pattern associated with fragmented mitochondria.

Fourth, to determine the functional state of the mitochondria in cells treated with caspofungin as described above, we incubated the cells with the mitochondrial vital dye, JC-1 (5mM; Molecular Probes, Inc.) for 15min at 30°C to assess their mitochondrial membrane potential ($\Delta\psi_m$). In healthy cells with high $\Delta\psi_m$, JC-1 forms aggregates that exhibit intense red fluorescence, while in apoptotic cells with low $\Delta\psi_m$, the dye remains in the monomeric form, which has green fluorescence [21-23]. Our experiments revealed that caspofungin-treated W303-1A cells had significantly more unhealthy cells with low $\Delta\psi_m$ ($30\pm 4\%$; $p < 0.05$; Figure 1D, H) than untreated controls ($0.3\pm 0.2\%$).

In toto, all of our data suggests that caspofungin induces PCD in *S. cerevisiae* cells. Notably, while our studies were in progress, a paper was published that reported that caspofungin induces PCD in the yeast, *Candida albicans*, accompanied with phenotypic markers characteristic of apoptosis and necrosis, including some of the markers described above [24]. We have repeated and confirmed these results suggesting that caspofungin kills with a similar

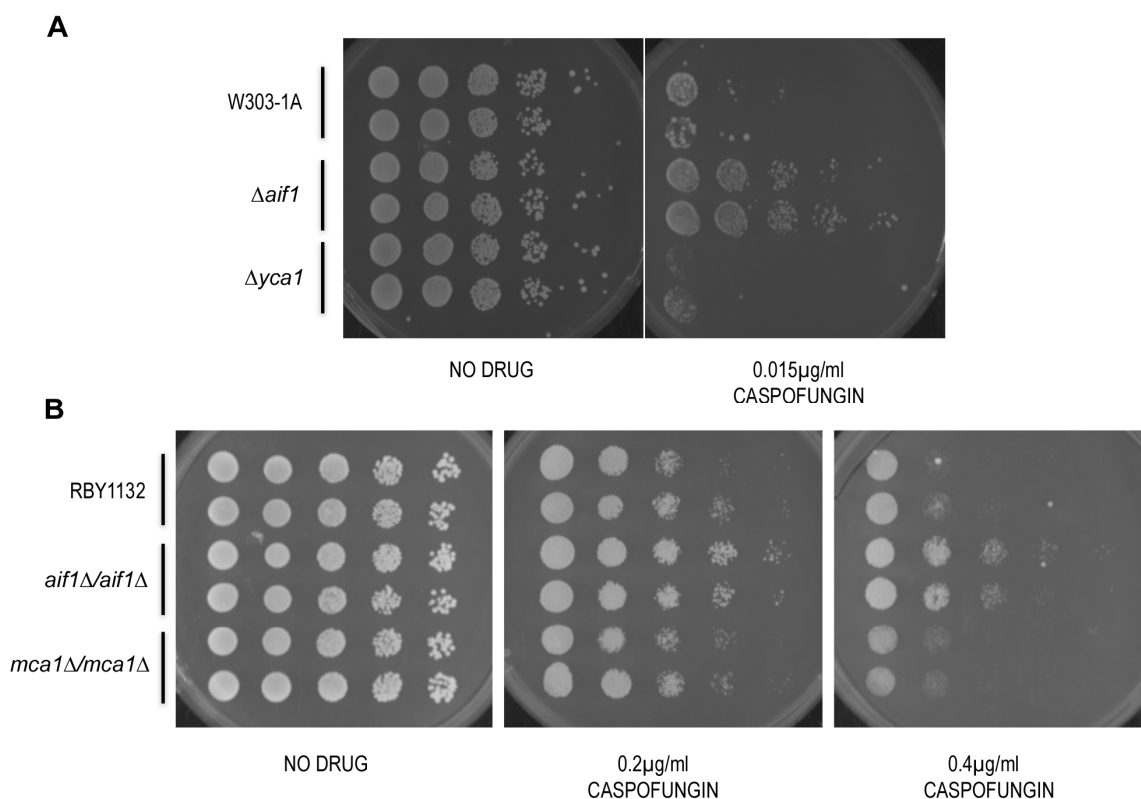


FIGURE 2: Deletion of AIF1 but not YCA1/MCA1 protects *Saccharomyces cerevisiae* and *Candida albicans* cells from caspofungin-induced programmed cell death. (A) A series of ten-fold dilutions of two cultures each of wildtype, $\Delta aif1$, and $\Delta yca1$ W303-1A *Saccharomyces* cells were plated on YPD plates and YPD plates with 0.015 $\mu\text{g/ml}$ caspofungin for two days. **(B)** A series of ten-fold dilutions of two independent cultures each of wildtype, $aif1\Delta/aif1\Delta$ and $mca1\Delta/mca1\Delta$ RBY1132 *Candida* cells were plated on YPD plates and YPD plates with the indicated concentrations of caspofungin for two days. Representative images are shown.

mechanism of action in both *Saccharomyces cerevisiae* and *Candida albicans* (data not shown).

Finally, ROS-mediated PCD in yeast is often distinguished into two different types depending upon their mechanism of killing, either caspase-dependent or caspase-independent yeast apoptosis [18, 25, 26]. Caspase-dependent cell death can be bypassed, at least in part, by the deletion of the yeast metacaspase, *YCA1/MCA1* [6, 27, 29]. In contrast, caspase-independent cell death cannot be suppressed by a *YCA1/MCA1* deletion [29]. Importantly, the yeast homolog of the mammalian apoptosis-inducing factor, called *AIF1* in both *Saccharomyces cerevisiae* and *Candida albicans*, has often been linked to caspase-independent cell death [7, 25].

While the recent paper showed that caspofungin induces PCD in *Candida*, it did not determine the mechanism of killing for the drug [24]. Indeed, the authors proposed that future studies should determine if either the caspase-dependent or the independent pathway is also involved in regulating yeast apoptosis upon exposure of caspofungin. To see if caspofungin induces either caspase-dependent or caspase-independent cell death, we spotted a series of ten-fold dilutions of two cultures each of wildtype, $\Delta aif1$, and $\Delta yca1$ W303-1A cells on YPD plates and YPD plates with caspofungin. After two days, we discovered that the viability of cells lacking the metacaspase was indistinguishable from that of wildtype cells (Figure 2A). This suggests that the caspase-like activity that we observed with our FLICA assay described above is associated with caspases and/or proteases other than Yca1p [18]. However, $\Delta aif1$ cells were relatively more resistant to caspofungin killing suggesting that this anti-fungal agent triggers *AIF1*-dependent PCD in *Saccharomyces cerevisiae* (Figure 2A). We repeated these experiments with wildtype, $aif1\Delta/aif1\Delta$, and $mca1\Delta/mca1\Delta$ mutants of *Candida albicans*, generously provided by Richard Bennett (Brown University). *MCA1* is the *Candida albicans* homolog of the *Saccharomyces cerevisiae* metacaspase gene, *YCA1* [16], while *AIF1* (orf19.3362) is the single homolog of the *Saccharomyces cerevisiae* *AIF1* gene [7, 25]. The assay was repeated with two independent isolates for each *Candida* null mutant to ensure that the observed phenotype is due to the knockout of the ORF and not to some spurious mutation. Once again, after two days incubation, we discovered that $aif1\Delta/aif1\Delta$ cells were more resistant to caspofungin than either wildtype or $mca1\Delta/mca1\Delta$ cells (Figure 2B).

In conclusion, our study suggests that caspofungin leads to caspase-independent cell death requiring *AIF1* in both *Saccharomyces cerevisiae* and *Candida albicans*. Of significance, though caspase-independent cell death has been observed in *Candida* before [29], this is the first time that *AIF1* has been implicated in this process for this pathogenic yeast. In sum, our findings reveal an additional dimension of this echinocandin's mechanism of action as an anti-fungal agent. They also point to a novel scenario for the acquisition of caspofungin resistance since this is the first time that drug resistance in *Candida* has been linked to a gene involved in cell death.

MATERIALS AND METHODS

Yeast Strains and Growth Conditions

All experiments were done with isogenic *Saccharomyces cerevisiae* strains in the W303-1A background (MATa *ade2*, *his3*, *leu2*, *trp1*, *ura3*, *ssd1-d2*). The $\Delta yca1$ mutant (SKY2880) was previously described [30]. The $\Delta aif1$ mutant was created by disrupting the ORF with the kanR marker using a PCR-based knock-out strategy [31] and verified both by PCR and by phenotypic analysis. Experiments with *Candida albicans* were done with wildtype, $aif1\Delta/aif1\Delta$, and $mca1\Delta/mca1\Delta$ mutants of *Candida albicans*, generously provided by Richard Bennett (Brown University). Mutants were created in the wildtype RBY1132 strain background (MTL a/a, *leu2/leu2 his1/his1 arg4/arg4*) by disrupting the ORF with a disruption cassette generated by fusion PCR and verified as previously described [32]. For all the experiments described in this paper, cells were cultured using standard protocols [33]. Caspofungin (*Candidas*) was generously provided by Merck (Rahway, NJ) and by Timothy Flanigan (Rhode Island Hospital/Warren Alpert School of Medicine at Brown University). Unless noted otherwise, all other drugs and reagents were purchased from SIGMA-Aldrich.

Caspofungin Activity Assay

Wildtype budding yeast cells in the W303-1A background were cultured in YPD overnight. In the morning, they were reinoculated in fresh media and allowed to grow for several hours until they reached the exponential phase (OD₆₀₀ 0.2-0.4). They were then resuspended at the same cell density in fresh media or in fresh media containing 0.02 µg/ml caspofungin for three hours, a drug concentration below the MIC of 0.04 µg/ml for this strain (data not shown). Next, the cells were washed and resuspended in a solution of YPD and 50 µg/ml propidium iodide. Propidium iodide is a membrane impermeable fluorescent molecule that is excluded from viable cells. Fluorescent cells were counted with either a Zeiss Axiophot Fluorescence Microscope or a Zeiss LSM 700 Laser Confocal Microscope fitted with Filter Cube 43 Cy3/Rhod/RFP (EX BP 545/25, BS FT 565, EM BP 606/70). Measurements were repeated with at least three independent cell cultures.

Apoptosis Assays

Wildtype budding yeast cells in the W303-1A background were cultured in YPD overnight. In the morning, they were reinoculated in fresh media and allowed to grow for several hours until they reached the exponential phase (OD₆₀₀ 0.2-0.4). They were then resuspended at the same cell density in fresh media or in fresh media containing 0.02 µg/ml caspofungin for three hours. To detect the presence of reactive oxygen species (ROS), they were stained with 2.5 µM CellROX Green (Molecular Probes, Inc.) for 10min at room temperature. This cell-permeable dye exhibits bright fluorescence upon oxidation by ROS, which was visualized with a Zeiss LSM700 confocal microscope fitted with Filter Cube 38 GFP (EX BP 470/40, BS FT 495, EM BP 525/50). Next, to determine the presence of activated caspase-like activity, they were stained with a FLICA apoptosis detection kit (ImmunoChemistry Technologies, LLC) according to the manufacturer's specifications, and imaged with a Zeiss LSM700 confocal microscope fitted with Filter Cube 38 GFP. Third, to determine the functional state of the mitochondria in cells treated with caspofungin as described above, they were incubated with the mitochondrial vital dye, JC-1 (5mM; Molecular Probes, Inc.) for 15min at 30°C to assess

their mitochondrial membrane potential ($\Delta\psi_m$). In healthy cells with high $\Delta\psi_m$, JC-1 forms aggregates that exhibit intense red fluorescence, while in apoptotic cells with low $\Delta\psi_m$, the dye remains in the monomeric form, which has green fluorescence [21-23]. Both were imaged with the Zeiss LSM700 fitted either with Filter Cube 43 Cy3/Rhod/RFP or with Filter Cube 38 GFP. Finally, to detect the fragmentation of mitochondria, the cells were transformed with a p416 GPD-mtGFP plasmid constitutively expressing a mitochondria-localized GFP marker [19], treated with caspofungin as described above, and imaged with a Zeiss LSM 700 Laser Confocal Microscope fitted with Filter Cube 38 GFP. For all the apoptosis assays, the experiment was repeated with at least three independent cultures. Representative images are shown with consistent parameters applied to comparable panels in each figure. Statistical significance was determined with the unpaired Student's *t*-test, and a minimum of 500 cells was counted for each assay.

Spot Assays

Wildtype and mutant cells of the indicated genotypes were grown overnight in YPD at 30°C (*Saccharomyces*) or at 37°C (*Candida*) respectively. For each yeast strain, a series of 10-fold dilutions was then prepared in water with cells in the exponential phase (OD₆₀₀ 0.2-0.4) over a range of concentrations from 10⁻¹ to 10⁻⁵ relative to the initial culture. Spots of 5 μ l from each dilution series were then plated on the indicated media and cultured at 30°C (*Saccharomyces*) or at 37°C (*Candida*) for 2 days. Plates supplemented with caspofungin were poured and used on the same day. All spot assays were repeated at least three times and a representative experiment is shown.

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CONFLICT OF INTEREST

The authors declare no conflict of interest.

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