

1 **Heightened efficacy of anidulafungin when used in combination with manogepix or 5-**
2 **flucytosine against *Candida auris* in vitro.**

3 Running title: Synergistic drug combinations against *Candida auris*.

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19

20 **Abstract**

21 *Candida auris* is an emerging, multi-drug resistant fungal pathogen that causes refractory
22 colonisation and life-threatening invasive nosocomial infections. The high proportion of *C. auris*
23 isolates that display antifungal resistance severely limits treatment options. Combination
24 therapies provide a possible strategy to enhance antifungal efficacy and prevent the emergence of
25 further resistance. Therefore, we examined drug combinations using antifungals that are already
26 in clinical use or undergoing clinical trials. Using checkerboard assays we screened combinations
27 of 5-flucytosine and manogepix (the active form of the novel antifungal drug fosmanogepix)
28 with anidulafungin, amphotericin B or voriconazole against drug resistant and susceptible *C.*
29 *auris* isolates from clades I and III. Fractional inhibitory concentration indices (FICI values) of
30 0.28-0.75 and 0.36-1.02 were observed for combinations of anidulafungin with manogepix or 5-
31 flucytosine, respectively, indicating synergistic activity. The high potency of these anidulafungin
32 combinations was confirmed using live-cell microfluidics-assisted imaging of fungal growth. In
33 summary, combinations of anidulafungin with manogepix or 5-flucytosine show great potential
34 against both resistant and susceptible *C. auris* isolates.

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36 Keywords: *Candida auris*; antifungal combination; anidulafungin; flucytosine; manogepix;
37 synergy

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41 **Introduction**

42 *Candida auris* is an emerging fungal pathogen that causes nosocomial invasive infections and
43 that is difficult to eradicate following colonisation of hospitalised patients (1). *C. auris* was first
44 identified in 2009 in Japan, but since then outbreaks have been observed on most continents (1,
45 2). *C. auris* strains have been subdivided into four genetic clades, the South Asian (I), East Asian
46 (II), South African (III) and South American (IV) clades (3), with a potential fifth Iranian clade
47 identified more recently (4). The organism colonises the skin and can lead to mucosal or
48 bloodstream infections, predominately in immunocompromised hosts (1). Invasive *C. auris*
49 infections are associated with mortality rates between 28% and 60%, and treatment failure due to
50 antifungal resistance is often observed (1, 3, 5–11).

51 To date, only four classes of antifungal drug are available for the treatment of invasive fungal
52 infections: the azoles, polyenes, echinocandins and the nucleoside analogue 5-flucytosine. 5-
53 flucytosine has high oral bioavailability with high activity against *C. auris*, but it is not generally
54 used in monotherapy due to the rapid emergence of resistance (12). Current guidelines
55 recommend echinocandin treatment as first line therapy for invasive candidiasis and for *C. auris*
56 infection in particular (13, 14). However, echinocandin resistance can develop during treatment
57 (15, 16). Resistance to all four existing classes of antifungal has been reported in *C. auris*, with
58 varying drug susceptibilities and resistance mechanisms between clades (17). Around 90 % of
59 *C. auris* isolates show resistance to fluconazole with varying susceptibilities to other azoles (3, 6,
60 9, 18). Resistance to amphotericin B and the echinocandins appears to be less common, having
61 been reported in 13-35 % and 2-7 % of tested isolates, respectively (3, 9, 18). Alarmingly,
62 between 3 % and 41 % of isolates exhibit resistance to two or more antifungal classes (3, 18).
63 Consequently, the Centers for Disease Control and Prevention (CDC) recently added *C. auris* to

64 its list of urgent antibiotic resistance threats (19) and the World Health Organisation (WHO)
65 declared it a critical threat in its fungal priority pathogens list (14).

66 The limited number of antifungal drugs as well as the increased threat of antifungal resistance in
67 *C. auris* means that novel treatment strategies are urgently needed. Combinations of antifungals
68 with different mechanisms of action provide one proposed therapeutic strategy. Previous *in vitro*
69 studies investigated combinations of echinocandins with azoles or the polyene amphotericin B
70 (20–24) and combinations of 5-flucytosine with the other three antifungal classes in *C. auris*
71 (25–27). These studies observed either synergy or indifference and no antagonism for all of the
72 tested combinations, with variability between *C. auris* isolates. The most promising
73 combinations were azoles combined with echinocandins which, in two studies, resulted in
74 synergy against all tested isolates (20, 23).

75 Combinations with 5-flucytosine are of particular interest as its combinations with amphotericin
76 B and fluconazole have been shown to be superior to monotherapy in phase III clinical trials
77 against cryptococcal meningitis (28). As a result of these trials, 5-flucytosine is now more widely
78 available globally, including in countries such as South Africa which suffers a high burden of
79 *C. auris* candidemia (28, 29). Echinocandin combinations with 5-flucytosine have been reported
80 to be indifferent in most cases, but these combinations have shown 100% growth inhibition and
81 fungicidal activity against multidrug-resistant isolates (25–27).

82 None of these studies included the new antifungal fosmanogepix, which has recently completed
83 phase 1 and 2 clinical trials, and is one of several new antifungals in the pipeline that may exhibit
84 activity also against *C. auris* (30). Fosmanogepix is a prodrug that is converted to the active
85 compound manogepix by systemic phosphatases (31). Manogepix inhibits a novel antifungal
86 target, Gwt1, which is involved in the GPI-anchor biosynthetic pathway, leading to a decrease in

87 cell wall-anchored mannoproteins (31). In the present study, we examined combinations of
88 manogepix or 5-flucytosine with anidulafungin, amphotericin B or voriconazole against a range
89 of resistant and susceptible *C. auris* isolates *in vitro*.

90 **Material and Methods**

91 *Fungal isolates*

92 Twenty-five clinical *C. auris* isolates belonging to clades I, III and IV isolated from 6 patients
93 from a range of sites (blood, urine, respiratory tract, skin) were obtained from the CDC (Table
94 1). Clade designations were based on whole genome sequencing (Gifford *et al.*, in preparation).
95 Isolates were maintained at - 80 °C in 25 % glycerol broth and subcultured on Sabouraud
96 dextrose agar (SDA) at 37 °C for up to 48 h.

97 *Antifungal susceptibility testing*

98 Antifungal susceptibility testing was performed using the broth microdilution method according
99 to EUCAST guidelines (32). Flat-bottom, tissue-treated 96-well plates were used. Anidulafungin
100 (MedChem Express), amphotericin B (Merck), fluconazole (Thermo Scientific), 5-flucytosine
101 (Thermo Scientific), fosmanogepix (MedChem Express), manogepix (MedChem Express) and
102 voriconazole (Sigma Aldrich) were dissolved in 100 % dimethyl sulfoxide (DMSO). The range
103 of antifungal concentrations tested were 0.016 to 8 mg/L for anidulafungin, 0.03 to 16 mg/L for
104 amphotericin B and voriconazole, 0.25 to 128 mg/L for fluconazole, 0.008 to 4 mg/L for 5-
105 flucytosine, 0.004 to 2 mg/L for fosmanogepix and 0.002 to 1 mg/L for manogepix. Antifungal
106 dilution series were prepared in RPMI supplemented with glucose to 2 % and buffered at pH 7
107 using 3-(N-morpholino) propanesulfonic acid (MOPS) at a final concentration of 0.165 mol/L
108 (RPMI 2%G-MOPS). Spectrophotometer readings at 530 nm were taken after incubation at
109 37 °C for 24 h The minimum inhibitory concentration (MIC) endpoint for amphotericin B was

110 defined as the lowest concentration leading to 90 % reduction in growth compared to the drug-
111 free control (MIC₉₀), while MIC₅₀ endpoints, measuring 50 % reduction in growth compared to
112 the drug-free control, were used for all other antifungal agents. Tentative CDC breakpoints for *C.*
113 *auris* were used to define resistance to anidulafungin (≥ 4 mg/L), amphotericin B (≥ 2 mg/L),
114 fluconazole (≥ 32 mg/L) and voriconazole (≥ 2 mg/L) ([https://www.cdc.gov/fungal/candida-](https://www.cdc.gov/fungal/candida-auris/c-auris-antifungal.html)
115 [auris/c-auris-antifungal.html](https://www.cdc.gov/fungal/candida-auris/c-auris-antifungal.html)). A known issue for broth microdilution susceptibility testing of
116 amphotericin B in RPMI medium is the clustering of MICs around the breakpoint of 2 mg/L
117 making it difficult to distinguish resistant and susceptible isolates (33). There are no breakpoints
118 available for 5-flucytosine and fosmanogepix. *Candida krusei* ATCC 6258 and *Candida*
119 *parapsilosis* ATCC 22019 were used as quality control strains as recommended by the EUCAST
120 guidelines (32). All experiments were performed in triplicate.

121 *Antifungal combination testing*

122 Interactions of antifungal drugs were tested using checkerboard assays based on EUCAST
123 guidelines (32). The range of antifungal concentrations tested was dependent on the MIC of each
124 isolate, with the highest concentration at 4 x MIC. Columns 3 to 12 of a 96-well microtiter plate
125 were filled with 50 μ l of drug A and rows B to H were filled with 50 μ l of drug B. Column 1
126 served as drug-free growth and sterility control. The inoculum was prepared by suspending five
127 distinct colonies from 40- 48h-old cultures in distilled water, counting the cell number using a
128 haemocytometer and adjusting inocula to 5×10^5 cells/ml. The plates were inoculated with 100
129 μ l and incubated at 37 °C for 24 h. OD readings were taken after 24 h using a spectrophotometer
130 at 530 nm. All experiments were performed in triplicate.

131 Two different approaches were applied in the analysis of drug interactions. The fractional
132 inhibitory concentration index (FICI) was calculated as follows:

$$FICI = \frac{C_A}{MIC_A} + \frac{C_B}{MIC_B}$$

133 C_A and C_B are the concentrations of the drugs A and B in combination and MIC_A and MIC_B are
134 the MICs of the drugs alone. MIC values were rounded to the next highest two-fold
135 concentration if the endpoint was not reached within the tested concentration range. The
136 interaction was considered synergistic for $FICI \leq 0.5$, partially synergistic between >0.5 and <1.0 ,
137 additive at 1.0, indifferent between >1.0 and <4 and antagonistic >4 (24). In the following, the
138 term “any synergy” refers to FICI values of <1 , thereby including complete and partial synergy.
139 In the presence of antagonism, the maximum median FICI values were reported, otherwise
140 minimum median FICI values were given. Additionally, drug interactions were visualised using
141 a response surface analysis approach with Combenefit software (version 2.021) under application
142 of the Bliss independence model (34).

143 *Microfluidics imaging*

144 *C. auris* B12663 cells were grown and prepared as described above. Inocula were adjusted to
145 2×10^5 cells/ml. Antifungal mono- and combination treatments were prepared in RPMI 2%G-
146 MOPS at the MIC. CellASIC® ONIX Y04C microfluidic plates (Millipore Merck) were washed
147 with RPMI 2%G-MOPS by applying 5 psi perfusion for 5 min using the CellASIC® ONIX2
148 microfluidic system (version 1.0.4 Millipore Merck). Yeasts were loaded into the CellASIC
149 culture chambers by applying 8 psi for 5 s twice (Thomson *et al.*, in preparation). Adhered cells
150 were then perfused with RPMI 2%G-MOPS for 4 h at 1 psi. After 4 h, cells were exposed to the
151 antifungal(s), or to RPMI 2%G-MOPS for the drug-free control, by applying 5 psi for 5 min,
152 followed by perfusion at 1 psi for 20 h at 37 °C, during which the microfluidic plates were
153 subjected to multi-point 4D imaging on an inverted AxioObserver Z1 microscope (Carl Zeiss).

154 Differential interference contrast (DIC) images were captured with a 20x/0.8NA
155 PlanApochromatic DIC objective and a 16-bit ORCA-Fusion sCMOS camera (Hamamatsu). The
156 area of colonies over time was measured in FIJI 1.53t (35) using an adapted method for
157 migration analysis from Venter and Niesler (36). Briefly, during the time series, colony edges
158 were found (Process → Find Edges), the image blurred fifteen times (Process → Smooth) and
159 inverted (Edit → Invert) before thresholding (Image → Adjust → Threshold: Default) to
160 quantify the total fungal area (Analyse → Analyse Particles). Increases in 2-dimensional colony
161 area were used to calculate the doubling times.

162 **Results**

163 *Antifungal activity against C. auris isolates*

164 The antifungal susceptibility profiles of 25 *C. auris* isolates were determined in order to select a
165 subset of isolates with different drug susceptibilities for antifungal combination testing. The
166 ranges of MIC values for the *C. auris* isolates against the tested antifungals are summarised in
167 Table 2 and Table S1. MIC₉₀ values for amphotericin B clustered around the breakpoint of 2
168 mg/L which is a known problem for broth microdilution susceptibility testing of amphotericin B
169 in RPMI medium, making it difficult to distinguish resistant and susceptible isolates (33).
170 Fluconazole showed a large percentage of resistant *C. auris* isolates (96 %; breakpoint
171 ≥32 mg/L) with high MIC₅₀ values ranging from 4 to ≥128 mg/L, while the other triazole tested
172 (voriconazole) displayed more potent antifungal activity with MIC₅₀ ranging from 0.06 to 16
173 mg/L and 40 % resistant isolates (breakpoint ≥2 mg/L). Of all the antifungals tested with an
174 available breakpoint, anidulafungin produced the lowest percentage of resistant isolates (32 %;
175 ≥4 mg/L). The most potent antifungal activity against *C. auris* was observed for manogepix

176 (MIC₅₀/MIC₉₀, 0.008/0.03 mg/L; range, 0.004-0.03) followed by 5-flucytosine (MIC₅₀/MIC₉₀,
177 0.25/0.25 mg/L; range, 0.125-0.25).

178 *Interaction of antifungal drug combinations against C. auris isolates*

179 Based on their MIC values, 11 *C. auris* isolates with different drug susceptibility profiles were
180 selected to investigate the interactions of anidulafungin, amphotericin B and voriconazole with 5-
181 flucytosine or manogepix. The FICI values for these combinations, as determined by the
182 checkerboard assays, are presented in Table 3 and Figure 1 (FICI values of separate repeats can
183 be found in Tables S2 and S3). The combination of anidulafungin with 5-flucytosine resulted in
184 synergistic interactions for 10/11 isolates (synergy, 2/11 isolates; partial synergy, 8/11 isolates).
185 Meanwhile the combination of anidulafungin with manogepix led to synergy in all 11 isolates
186 (synergy, 5/11 isolates; partial synergy, 6/11 isolates). These FICI values corresponded to a
187 median (range) decrease in MIC₅₀ of 2 log₂-fold (1- to 4 log₂-fold) for anidulafungin and 2 log₂-
188 fold (0- to 4 log₂-fold) for 5-flucytosine (Figure 2A), or 3 log₂-fold (1- to 9 log₂-fold) for
189 anidulafungin and 2 log₂-fold (1- to 3 log₂-fold) for manogepix (Figure 2B). Additionally, both
190 anidulafungin combinations achieved fungistatic activity with a log₁₀-fold reductions in CFUs/ml
191 of 2.2 and 0.8 compared to the starting inoculum for the combination with manogepix and 5-
192 flucytosine, respectively, while the corresponding monotherapies only had a negligible
193 antifungal effect (Figure S7).

194 The combination of amphotericin B with 5-flucytosine did not show full synergy for any of the
195 tested isolates, though partial synergy was observed in 4/11 isolates (median FICIs 0.63-0.75).
196 The other isolates showed either additive (5/11 isolates) or indifferent (2/11 isolates, median
197 FICIs 1.01) interactions for amphotericin B with 5-flucytosine. For the combination of
198 manogepix and 5-flucytosine, 3/11 isolates displayed partial synergy (median FICIs 0.54-0.58)

199 and 4/11 isolates showed additive or indifferent interactions (median FICIs 1.01). The
200 combination of manogepix and 5-flucytosine led to large reductions in the MIC₅₀ by median
201 (range) 7 log₂-fold (1- to 8 log₂-fold) for 5-flucytosine, while the manogepix MIC₅₀ were only
202 decreased by median (range) 0 log₂-fold (0- to 2 log₂-fold) (Figure S1C). The drug combination
203 resulting in the least favourable interactions was voriconazole with 5-flucytosine with 3/11
204 isolates displaying antagonistic interactions (median FICIs 4.48-4.50), and the remaining isolates
205 displaying additive (3/11 isolates) or indifferent (5/11 isolates, median FICIs 1.01) interactions.

206 Response surface analyses were also used to examine the drug combinations, and an example is
207 shown in Figure 3 for the multidrug-resistant isolate B12663 (see Figures S2-S6 for the other
208 isolates). Consistent with the FICI scores, the synergy maps indicate synergy for the combination
209 of anidulafungin and manogepix (median FICI 0.33) and weak synergy for combinations of 5-
210 flucytosine with anidulafungin (median FICI 0.74) or amphotericin B (median FICI 0.75). In
211 contrast to the FICI calculation, which only focuses on drug concentrations corresponding to
212 MIC values, the response surface analysis permits the examination of drug interactions over a
213 wide range of tested concentrations. This revealed antagonism at the lower end of some
214 concentration ranges that was missed by the FICI approach, highlighting the concentration-
215 dependence of the interactions.

216 *Real time imaging of anidulafungin combinations against a multidrug-resistant C. auris isolate*
217 *using microfluidics*

218 A microfluidics imaging approach was employed to further investigate the effects, at a single-
219 cell level, of the two most promising drug combinations: anidulafungin with manogepix, and
220 anidulafungin with 5-flucytosine. This system is less static than the traditional microbroth
221 dilution method as the cells are constantly perfused with fresh medium containing different

222 antifungal drugs. Again, the multidrug-resistant *C. auris* isolate B12663 was chosen for analysis.
223 Both drug combinations showed dramatic effects upon cell growth, markedly reducing the size
224 of colonies compared to the relevant monotherapies and media-only controls (Figure 4A; Movies
225 S1 and S2). Doubling times, measured by 2-dimensional colony area changes, increased
226 significantly in the presence of the drug combinations compared to the individual antifungals. An
227 increase from 3.19 h (5-flucytosine alone) to 4.90 h ($p < 0.001$) was observed for anidulafungin
228 combined with 5-flucytosine (Figure 4B). Similarly, an increase from 2.75 h (manogepix alone)
229 to 9.50 h ($p < 0.001$) was seen for the anidulafungin-manogepix combination (Figure 4C). These
230 changes in doubling time correspond to 63.5 % (anidulafungin-5-flucytosine) and 96.5%
231 (anidulafungin-manogepix) decrease in colony area after 24 h compared to 5-flucytosine and
232 manogepix, respectively (data not shown). These findings were again consistent with those of the
233 checkerboard and response surface analysis experiments, in that the combination of
234 anidulafungin and manogepix showed the most potent impacts on cell growth, followed by the
235 combination of anidulafungin plus 5-flucytosine.

236 The cellular morphology was further examined at higher magnification after exposing the *C.*
237 *auris* cells to the antifungals in monotherapy or combination for 24 h (Figure S8). In drug-free
238 medium the cells had a well-defined, oval morphology. Under exposure to anidulafungin,
239 manogepix and both anidulafungin combinations the cells displayed a rounder morphology with
240 the formation of aggregates, while 5-flucytosine treatment resulted in a more elongated
241 phenotype. Additionally, enlarged, round cells were observed in the presence of manogepix and
242 both combinations.

243 **Discussion**

244 The emergence and global spread of multidrug-resistant *C. auris* strains poses a serious health
245 threat. The high prevalence of antifungal resistance reported for *C. auris* isolates (3, 6–9, 11, 18,
246 24) was also observed in the isolates used in this study, with the majority of isolates resistant to
247 fluconazole, 40 % resistant to voriconazole and 32 % resistant to anidulafungin. The ability of
248 *C. auris* to develop resistance to all of the available classes of antifungal drug severely limits
249 treatment options.

250 New antifungal drugs, such as fosmanogepix, are currently in development (reviewed in (30)).
251 *C. auris* currently appears susceptible to the active version of this new class of drugs
252 (manogepix), but there is a high risk of resistance developing following its introduction to the
253 clinic unless precautionary measures are taken. Combination therapies provide a proven strategy
254 that has already been employed in the treatment of viral and bacterial infections to prevent the
255 emergence of resistance to a single drug (37). Additionally, combination therapies have the
256 potential to improve efficacy through additive or synergistic interactions, allowing lower drug
257 doses to be used, thereby reducing dose-related toxicity.

258 Thus far, nine studies have examined antifungal drug combinations against *C. auris*. The
259 majority of these studies focussed on combinations of azoles with echinocandins (20, 23, 24, 38),
260 while a smaller number have evaluated polyene-echinocandin interactions (21, 22) or
261 combinations with 5-flucytosine (25–27). These studies reported mainly synergistic (including
262 partial synergy) or indifferent interactions, with inter-strain variability observed for some
263 combinations. None of these studies included manogepix. Both manogepix and 5-flucytosine
264 have potent antifungal activity against *C. auris* as shown here and observed by others (39–44).
265 Therefore, we examined interactions of the echinocandin anidulafungin, the azole voriconazole

266 and the polyene amphotericin B with either 5-flucytosine or manogepix using checkerboard
267 assays, response surface analyses and microfluidics imaging.

268 According to the FICI values and response-surface analyses, the most potent combination (with
269 respect to the number of *C. auris* isolates that displayed synergy) was anidulafungin plus
270 manogepix, followed by the combination of anidulafungin with 5-flucytosine. The high efficacy
271 of these combinations was also confirmed by microfluidics imaging, which revealed dramatic
272 reductions in fungal growth compared to the relevant monotherapies. The interactions between 5-
273 flucytosine with either amphotericin B or manogepix were additive or indifferent for the majority
274 of the isolates, while the combination of voriconazole with 5-flucytosine was indifferent or
275 antagonistic.

276 Applying our FICI thresholds, Bidaud and co-workers also reported mainly partially synergistic
277 or additive interactions for combinations of amphotericin B, voriconazole or micafungin with 5-
278 flucytosine (25). However, they did not observe the antagonism for the combination of
279 voriconazole with 5-flucytosine that we observed here. Another study reported 100 % growth
280 inhibition of amphotericin B or anidulafungin-resistant *C. auris* isolates for amphotericin B-5-
281 flucytosine combinations (0.25/1 mg/L) or anidulafungin-5-flucytosine combinations (0.008/1
282 mg/L) (26). Based on our OD₅₃₀ measurements, more than 90 % growth inhibition was also
283 achieved for the majority of susceptible and resistant isolates we analysed, and this growth
284 inhibition could be reached at lower concentrations for some isolates. To the best of our
285 knowledge, antifungal combinations with fosmanogepix/manogepix have not been studied
286 previously against *Candida* species. One recent study compared amphotericin B monotherapy
287 with the combination therapy of fosmanogepix and amphotericin B in invasive mouse infection
288 models of *Aspergillus fumigatus*, *Rhizopus arrhizus* var. *delemar* and *Fusarium solani* (45). In

289 all three models, mortality and fungal burden were significantly reduced in the mice treated with
290 the combination therapy compared to amphotericin B or fosmanogepix alone (45).

291 For the majority of combinations and isolates we examined, the interactions were partially
292 synergistic or additive. However, even these interactions could be of interest clinically, as the
293 ultimate goal is to reduce fungal burden with a view to supporting the immune system in clearing
294 the infection. This reduction in fungal growth could be clearly observed in the microfluidics
295 imaging for the combination of anidulafungin with 5-flucytosine, which only displayed a
296 partially synergistic interaction for the imaged isolate in the checkerboard assays. Furthermore,
297 partially synergistic or additive interactions can lead to reductions in the MICs, potentially
298 allowing for a lowering of antifungal doses, thereby reducing toxicity. Reductions in MICs for
299 partially synergistic, additive and indifferent combinations have also been observed by others
300 (20, 24) and Caballero and colleagues reported that additive combinations of isavuconazole-
301 echinocandin combinations against *C. auris* can result in fungistatic effects which were absent
302 for single agents in time-kill assays (23). This is similar to our results showing negligible
303 antifungal activity for anidulafungin, manogepix and 5-flucytosine in monotherapy, whereas the
304 combinations of these two antifungals with anidulafungin showed heightened efficacy with the
305 reductions in CFUs/ml approaching the cidal threshold. The lack of fungicidal activity of the
306 echinocandins against *C. auris* in time-kill assays has also been observed by others reporting
307 either a fungistatic effect or the complete absence of antifungal activity (22, 23, 46, 47). In
308 comparison to anidulafungin monotherapy, the anidulafungin combinations resulted in 2.1 and
309 3.6 log₁₀-fold reductions in CFUs/ml for 5-flucytosine and manogepix combinations,
310 respectively, highlighting their advantage over monotherapy.

311 Cost and additional toxicities are potential barriers to implementation of antifungal
312 combinations, and, to date, routine use of antifungal combinations has been largely confined to
313 cryptococcal infection. However, affordable generic echinocandins and 5-flucytosine are now
314 available, and short courses of 5-flucytosine are known to be very safe, giving feasible current
315 options to try to prevent the inevitable increase in *C. auris* resistance consequent on continued
316 use of monotherapies. Furthermore, early studies of combination approaches with new agents
317 such as fosmanogepix could expand the options for clinical evaluation and prolong their clinical
318 efficacy.

319 The synergistic interactions we observed for anidulafungin combined with manogepix or 5-
320 flucytosine were within clinically relevant concentrations in most cases. Serum anidulafungin
321 concentrations of up to 7 mg/L are achievable in patients (48, 49) which is above the
322 anidulafungin concentrations corresponding to synergistic interactions for most isolates. For 5-
323 flucytosine all concentrations we tested fall well below the achievable serum concentrations (48).
324 In the case of fosmanogepix, no clinical pharmacokinetics data is publicly available to our
325 knowledge. Several safety and pharmacokinetics clinical studies for fosmanogepix have been
326 completed, but no results are available yet (NCT02956499, NCT02957929, NCT03333005).
327 However, the manogepix concentrations at which synergy was observed were relatively low,
328 ranging between 0.002 and 0.03 mg/L.

329 It should be noted that the current study employed a relatively small number of isolates, and
330 there was an unequal representation of *C. auris* clades. Additionally, the clustering of
331 amphotericin B MIC₉₀ around the breakpoint made it difficult to categorise the isolates according
332 to their amphotericin B susceptibility. Hence, other susceptibility testing methods such as the
333 Etest are recommended (17).

334 In summary, combinations of anidulafungin with manogepix or 5-flucytosine show the highest
335 potential against the tested *C. auris* isolates. Further studies are needed to determine the
336 mechanisms that underlie these drug interactions and to evaluate their efficacy and safety in the
337 murine model and whether these combinations also protect against the development of
338 resistance.

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353 Resistance Award [MRF-160-0009-ELP-BICA-C0802].

354 **Conflicts of Interest**

355 EB and DDT were funded by Gilead Sciences. DDT received consultancy fee from OwlStone
356 Medical in the last 5 years. TB has received research funding from Gilead Sciences, MSD and

357 Pfizer; speaker fees from Gilead Sciences and Pfizer and Advisory Board fees from Gilead
358 Sciences and Mundipharma. MH reports grants and research funding from Astellas, Gilead,
359 MSD, Pfizer, Euroimmun, F2G, Pulmocide, IMMY, Mundipharma and Scynexis, outside the
360 submitted work. TSH has received speaker fees from Gilead and Pfizer, an investigator award to
361 institution from Gilead, and has served as advisor to F2G.

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543

544 **Figure and Table Legends**

545 Table 1. *Candida auris* isolates.

546 (50, 51)

547

548 Table 2. Antifungal MIC distribution for 25 *C. auris* isolates.

549

550 Table 1. FICI values for 5 antifungal combinations against eleven *C. auris* isolates.

551

552 Figure 1. *In vitro* interactions of AFG, MGX, AMB, VRC and 5FC according to the FICI values
553 for 11 *C. auris* isolates.

554 Minimum FICI values shown in absence of antagonism, otherwise maximum FICI values
555 reported. Drug interaction ranges are indicated by background colour: Synergy, dark green;
556 partial synergy, light green; indifference, white; antagonism, red. Symbols represent FICI values
557 of three independent experiments. 5FC, 5-flucytosine; AFG, anidulafungin; AMB, amphotericin
558 B; MGX, manogepix; VRC, voriconazole.

559

560 Figure 2. Changes in MIC values due to antifungal combinations for 11 *C. auris* isolates.
561 MIC values for 11 *C. auris* isolates in combinations of anidulafungin with 5-flucytosine (A) and
562 manogepix (B) compared to the antifungals in monotherapy as determined by checkerboard
563 assays. Symbols represent median values of three independent experiments. 5FC, 5-flucytosine;
564 AFG, anidulafungin; MGX, manogepix.

565

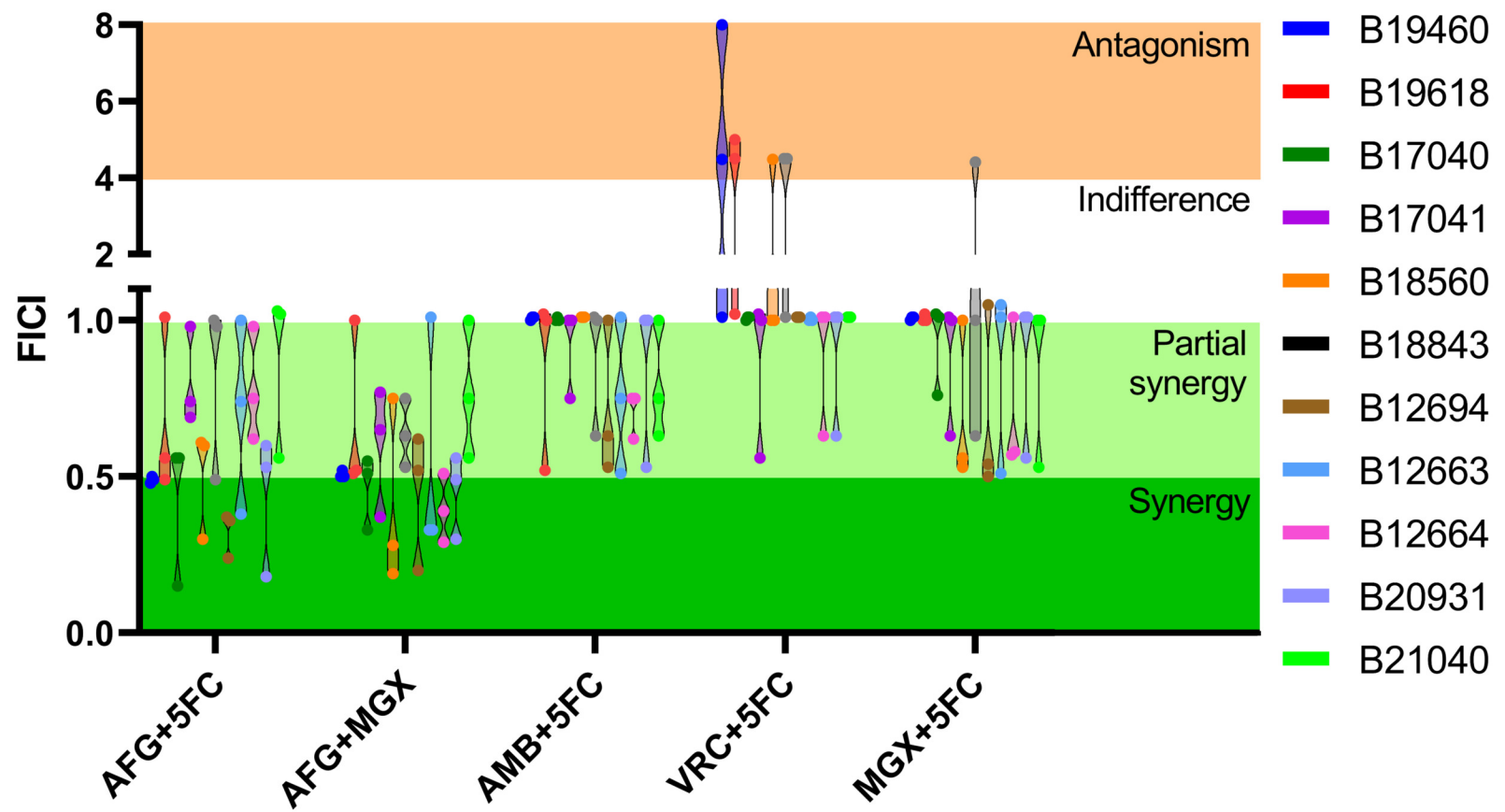
566 Figure 3. Synergy maps for 5 antifungal combinations against the multidrug-resistant *C. auris*
567 isolate B12663.

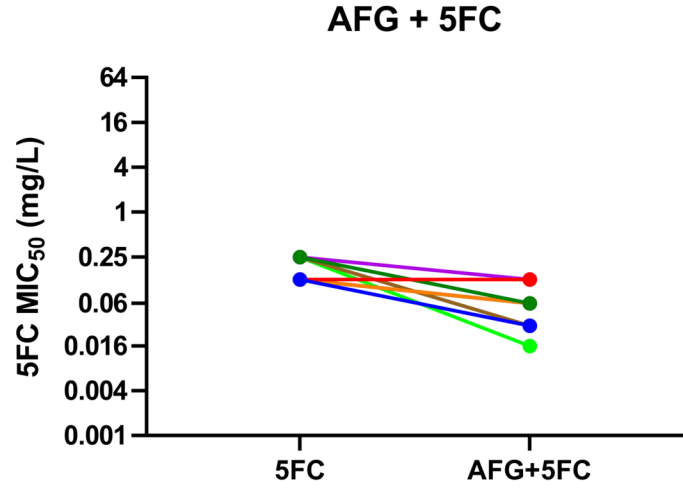
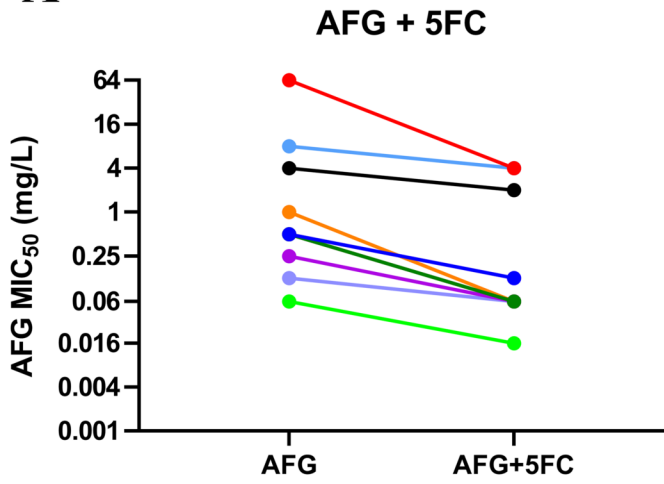
568 The interactions of 5-flucytosine with anidulafungin (A), amphotericin B (C) or voriconazole (D)
569 and the interactions of manogepix with anidulafungin (B) or 5-flucytosine (E) were analysed
570 with Combeneft (n=3). The graphs show the growth percentage relative to the drug-free control
571 with the colour scale representing the drug interaction. 5FC, 5-flucytosine; AFG, anidulafungin;
572 AMB, amphotericin B; MGX, manogepix; VRC, voriconazole.

573

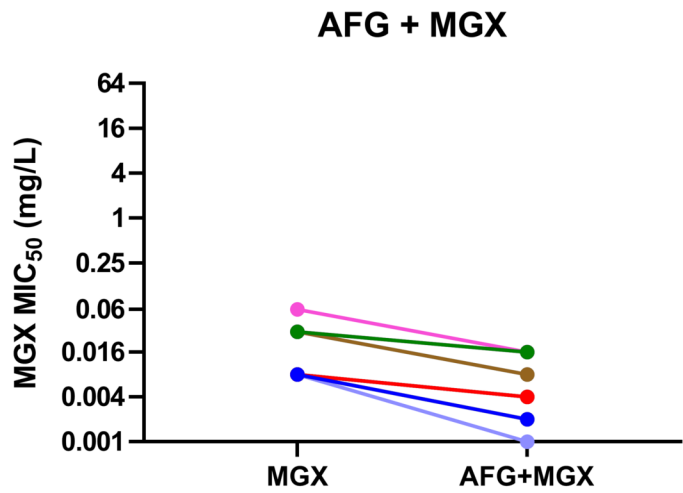
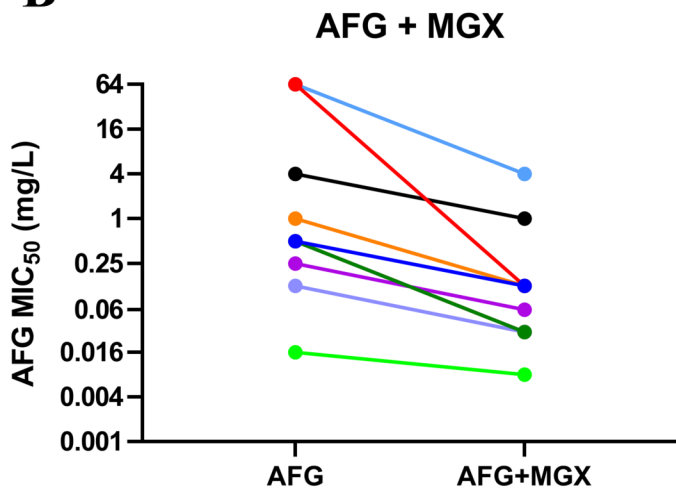
574 Figure 4. Microfluidics imaging of *C. auris* under antifungal combination exposure.

575 DIC images from two representative experiments (A) and doubling times (B, C) of *C. auris*
576 B12663 cells grown in the presence of RPMI 2%G-MOPS for 4 h, followed by further RPMI
577 2%G-MOPS or treatment with anidulafungin, 5-flucytosine and manogepix alone or in
578 combination at their MICs for 16 h. Doubling times were calculated by 2-dimensional colony
579 area changes for several colonies from two independent experiments. Mean \pm range. Scale bars:
580 100 μ m. *P \leq 0.05; **P \leq 0.01; ***P<0.001 (one-way ANOVA test with Bonferroni's correction).
581 5FC, 5-flucytosine; AFG, anidulafungin; MGX, manogepix.

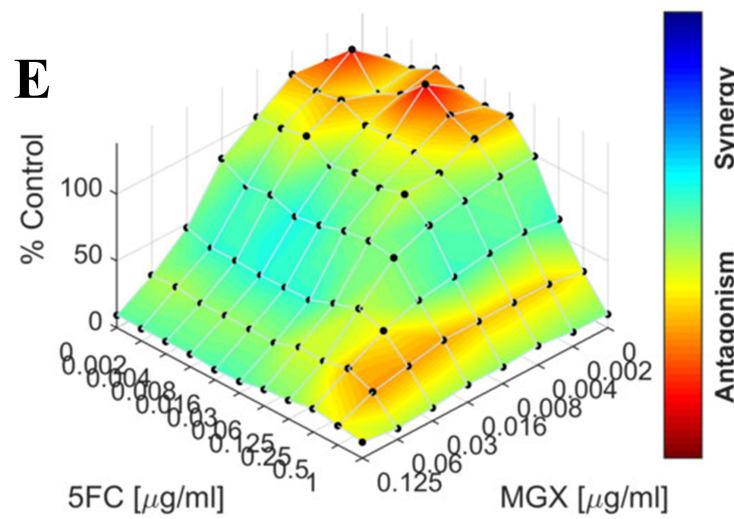
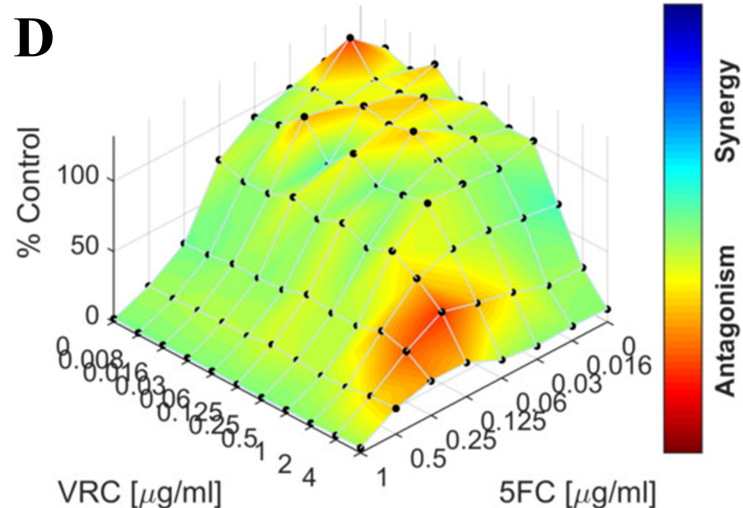
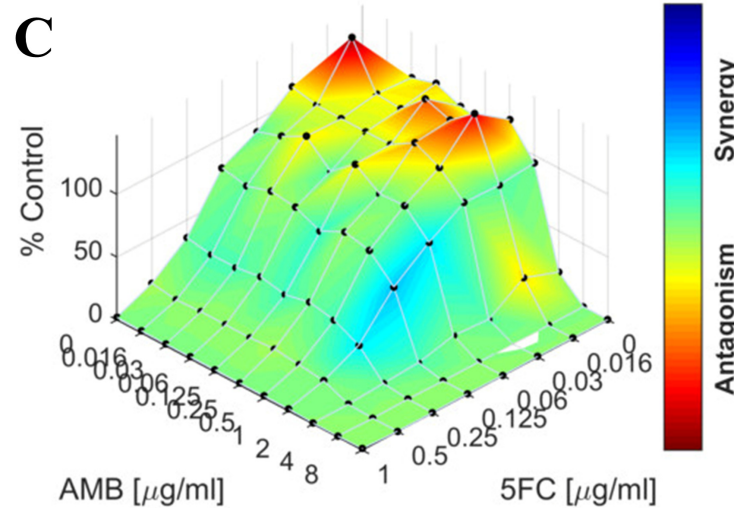
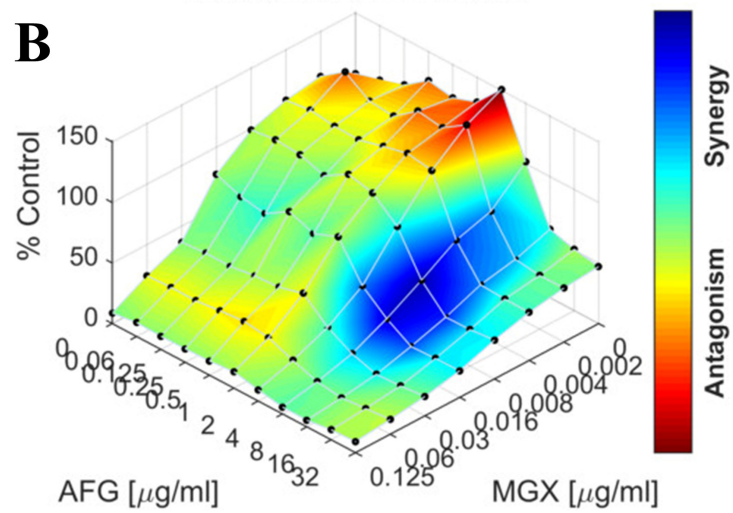
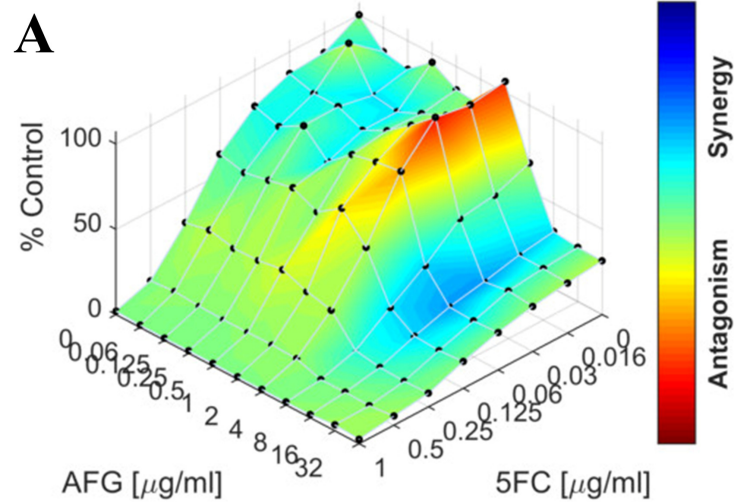


A

- B19460
- B19618
- B17040
- B17041
- B18560
- B18843
- B12694
- B12663
- B12664
- B20931
- B21040

B

- B19460
- B19618
- B17040
- B17041
- B18560
- B18843
- B12694
- B12663
- B12664
- B20931
- B21040



B12663 MIC values:

5FC: 0.25 mg/L

AFG: 8 mg/L

AMB: 2 mg/L

MGX: 0.03 mg/L

VRC: 2 mg/L

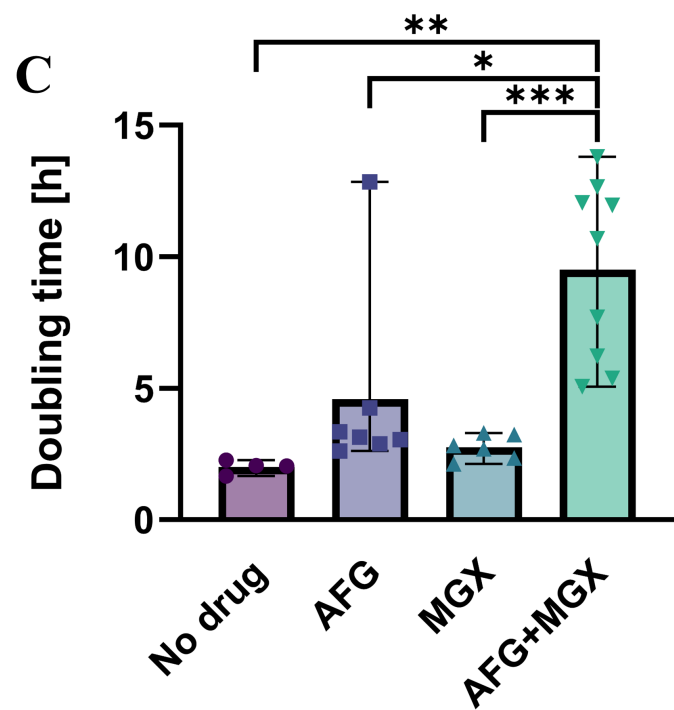
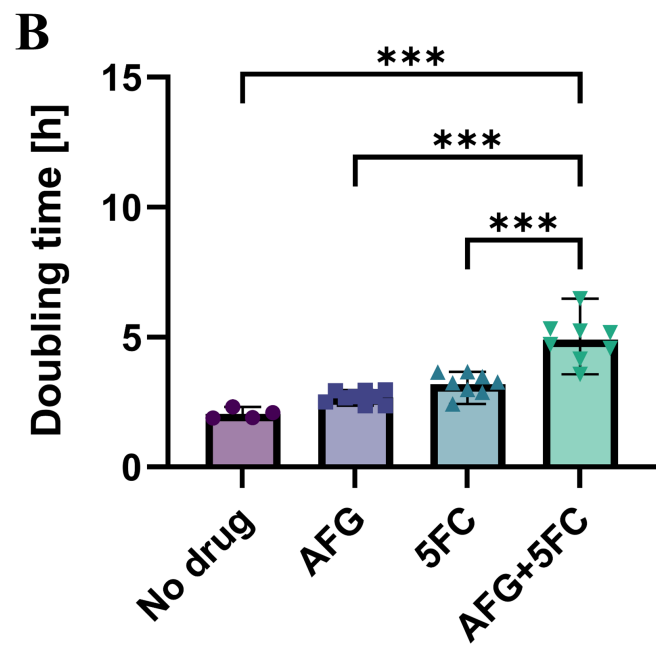
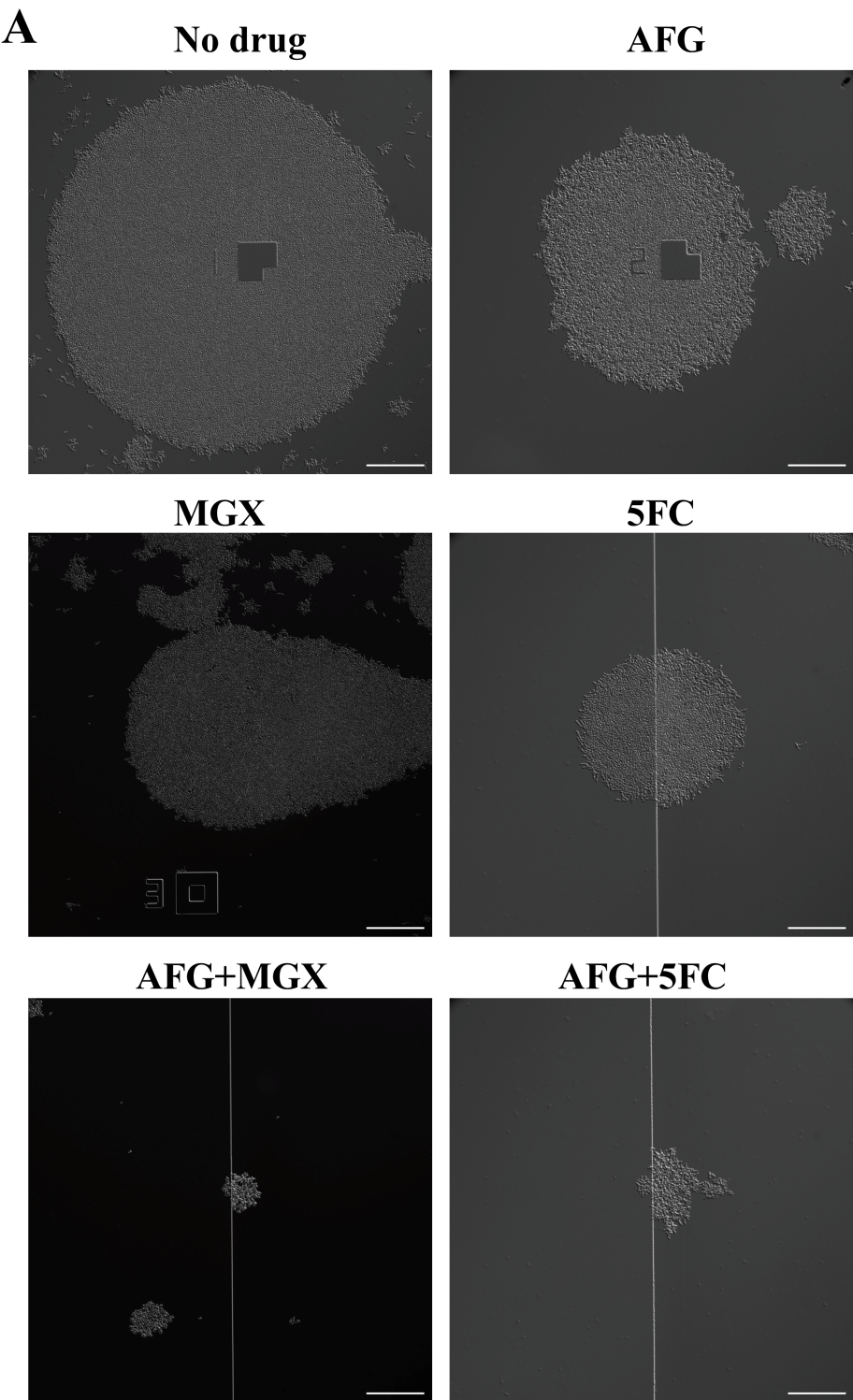


Table 1. *Candida auris* isolates.

Isolate number	Clade^a	Origin	Isolation day^b	Isolated from	Reference
B12406	South American	USA	Day 0	Patient A, Urine	(Chow, 2018)
B15223	South American	USA	Day 294	Patient A, Blood	
B19460	South Asian	USA	Day 0	Patient B, Sputum	
B19547	South Asian	USA	Day 16	Patient B, Unknown	
B19617	South Asian	USA	Day 46	Patient B, Urine	
B19837	South Asian	USA	Day 79	Patient B, Urine	
B19618	South Asian	USA	Day 62	Patient B, Urine	
B17040	South Asian	USA	Day 0	Patient C, Urine	
B17041	South Asian	USA	Day 15	Patient C, Sputum	
B17073	South Asian	USA	Day 44	Patient C, Urine	
B17201	South Asian	USA	Day 67	Patient C, Urine	
B18560	South Asian	USA	Day 0	Patient D, Blood	
B18845	South Asian	USA	Day 72	Patient D, Blood	
B18841	South Asian	USA	Day 103	Patient D, Blood	
B18843	South Asian	USA	Day 96	Patient D, Blood	
B12692	South Asian	USA	Day 11	Patient E, Rectal	(Di Pilato, 2021)
B12694	South Asian	USA	Day 0	Patient E, Groin swab	(Di Pilato, 2021)
B12663	South Asian	USA	Day 11	Patient E, Urine	(Di Pilato, 2021)
B12664	South Asian	USA	Day 11	Patient E, Respiratory	(Di Pilato, 2021)
B12688	South Asian	USA	Day 11	Patient E, Groin swab	(Di Pilato, 2021)
B20931	South African	USA	Day 0	Patient F, Blood	
B21040	South African	USA	Day 3	Patient F, Trachea Aspirate	
B21041	South African	USA	Day 3	Patient F, Groin swab	
B21042	South African	USA	Day 3	Patient F, Blood	
B21043	South African	USA	Day 3	Patient F, Blood	

^a Clade designation based on whole genome sequencing (Gifford *et al.*, in preparation).

^b In reference to isolation date of first isolate from respective patient.

Table 2. Antifungal MIC distribution for 25 *C. auris* isolates.

Drug	MIC (mg/L)														MIC ₅₀ ^a	MIC ₉₀ ^b	%R ^c			
	0.002	0.004	0.008	0.016	0.03	0.06	0.125	0.25	0.5	1	2	4	8	16				32	64	128
AMB					0	0	0	0	0	1	<u>24</u> ^d	0	0	0				2	2	96.0
FLC								0	0	0	0	1	0	0	5	5	<u>14</u>	≥128	≥128	96.0
VRC					0	1	2	6	1	5	<u>9</u>	0	0	1				1	2	40.0
AFG				0	3	3	5	3	2	0	1	0	<u>8</u>					0.25	≥8	32.0
5FC			0	0	0	0	11	<u>14</u>	0	0	0	0						0.25	0.25	No BP
MGX	0	<u>11</u>	5	1	8	0	0	0	0	0								0.008	0.03	No BP

^aMIC at which 50% of isolates were inhibited.

^bMIC at which 90% of isolates were inhibited.

^cPercentage of resistant isolates.

^dModal MICs are indicated with underlined numbers.

Grey background indicates tentative *C. auris* breakpoints according to the CDC.

5FC, 5-flucytosine; AFG, anidulafungin; AMB, amphotericin B; BP, breakpoint; FLC, fluconazole; MGX, manogepix; VRC, voriconazole.

Table 3. FICI values for 5 antifungal combinations against 11 *C. auris* isolates.

Isolate	AFG+5FC	AFG+MGX	AMB+5FC	VRC+5FC	MGX+5FC
	Median (range)	Median (range)	Median (range)	Median (range)	Median (range)
B19460	0.49 (0.48-0.50)	0.50 (0.50-0.52)	1.01 (1.00-1.01)	4.48 (1.01-8.00)	1.01 (1.00-1.01)
B19618	<u>0.56</u> (0.49-1.01)	<u>0.52</u> (0.51-1.00)	1.00 (0.52-1.02)	4.50 (1.02-5.00)	1.00 (1.00-1.02)
B17040	0.56 (0.15-0.56)	0.51 (0.33-0.55)	1.00 (1.00-1.01)	<u>1.01</u> (1.00-1.01)	1.01 (0.76-1.02)
B17041	0.74 (0.69-0.98)	0.65 (0.37-0.77)	1.00 (0.75-1.00)	<u>1.00</u> (0.56-1.02)	1.00 (0.63-1.01)
B18560	0.60 (0.30-0.61)	0.28 (0.19-0.75)	1.01 (1.01)	1.00 (1.00-4.48)	0.56 (0.53-1.00)
B18843	0.98 (0.49-1.00)	0.63 (0.53-0.75)	1.00 (0.63-1.01)	4.50 (1.01-4.50)	1.00 (0.63-4.41)
B12694	0.36 (0.24-0.37)	0.52 (0.20-0.62)	0.63 (0.53-1.00)	<u>1.01</u> (1.01)	0.54 (0.50-1.05)
B12663	<u>0.74</u> (0.38-1.00)	<u>0.33</u> (0.33-1.01)	0.75 (0.51-1.01)	<u>1.00</u> (1.00-1.01)	1.01 (0.51-1.05)
B12664	0.75 (0.62-0.98)	0.39 (0.29-0.51)	0.75 (0.62-0.75)	<u>1.01</u> (0.63-1.01)	0.58 (0.57-1.01)
B20931	0.53 (0.18-0.60)	0.49 (0.30-0.56)	1.00 (0.53-1.00)	1.01 (0.63-1.01)	1.01 (0.56-1.01)
B21040	1.02 (0.56-1.03)	0.75 (0.56-1.00)	0.75 (0.63-1.00)	1.01 (1.01)	1.00 (0.53-1.00)

Synergy, dark green; partial synergy, light green; indifference/additivity, white;

antagonism; red. Underlined values indicate resistance to either AFG or VRC.

5FC, 5-flucytosine; AFG, anidulafungin; AMB, amphotericin B; MGX, manogepix;

VRC, voriconazole