

1 ***In vitro* and *in vivo* effect of exogenous farnesol exposure against**
2 ***Candida auris***

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5 Running title: Farnesol against *Candida auris*
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9 Fruzsina Nagy^{1,2}, Eszter Vitális^{2,3}, Ágnes Jakab⁴, Andrew M. Borman⁵, Lajos Forgács^{1,2},
10 Zoltán Tóth^{1,2}, László Majoros¹, Renátó Kovács^{1,6*}
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13
14 ¹Department of Medical Microbiology, Faculty of Medicine, University of Debrecen,
15 Hungary

16 ²Doctoral School of Pharmaceutical Sciences, University of Debrecen, Hungary

17 ³Hospital Hygiene Ward, Clinical Centre, University of Debrecen, Hungary

18 ⁴Department of Molecular Biotechnology and Microbiology, Institute of Biotechnology,
19 Faculty of Science and Technology, University of Debrecen, Debrecen, Hungary.

20 ⁵Public Health England UK National Mycology Reference Laboratory, Bristol, United
21 Kingdom.

22 ⁶Faculty of Pharmacy, University of Debrecen, Hungary
23
24
25
26
27

28 *Corresponding author: Renátó Kovács, Department of Medical Microbiology, Faculty of
29 Medicine, University of Debrecen, 4032 Debrecen, Nagyerdei krt. 98., Hungary

30 Phone: 00-36-52-255-425, Fax: 00-36-52-255-424;

31 e-mail: kovacs.renato@med.unideb.hu
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43 **Abstract**

44 The spreading of multidrug-resistant *Candida auris* is considered as an emerging global
45 health threat. The number of effective therapeutic regimens is strongly limited; therefore,
46 development of novel strategies is needed. Farnesol is a quorum-sensing molecule with a
47 potential antifungal and/or adjuvant effect; it may be a promising candidate in alternative
48 treatment against *Candida* species including *C. auris*. To examine the effect of farnesol on *C.*
49 *auris*, we performed experiments focusing on growth, biofilm production ability, production
50 of enzymes related to oxidative stress, triazole susceptibility and virulence. Concentrations
51 ranging from 100 to 300 μM farnesol caused a significant growth inhibition against *C. auris*
52 planktonic cells for 24 hours ($p < 0.01-0.05$). Farnesol treatment showed a concentration
53 dependent inhibition in terms of biofilm forming ability of *C. auris*; however, it did not
54 inhibit significantly the biofilm development at 24 hours. Nevertheless, the metabolic activity
55 of adhered farnesol pre-exposed cells (75 μM) was significantly diminished at 24 hours
56 depending on farnesol treatment during biofilm formation ($p < 0.001-0.05$). Moreover, 300 μM
57 farnesol exerted a marked decrease in metabolic activity against one-day-old biofilms
58 between 2 and 24 hours ($p < 0.001$). Farnesol increased the production of reactive species
59 remarkably, as revealed by 2',7'-dichlorofluorescein (DCF) assay (3.96 ± 0.89 [nmol DCF
60 (OD_{640})⁻¹] and 23.54 ± 4.51 [nmol DCF (OD_{640})⁻¹] for untreated cells and farnesol exposed
61 cells, respectively; $p < 0.001$). This was in line with increased superoxide dismutase level
62 (85.69 ± 5.42 [munit (mg protein)⁻¹] and 170.11 ± 17.37 [munit (mg protein)⁻¹] for untreated
63 cells and farnesol exposed cells, respectively; $p < 0.001$), but the catalase level remained
64 statistically comparable between treated and untreated cells ($p > 0.05$). Concerning virulence-
65 related enzymes, exposure to 75 μM farnesol did not influence phospholipase or aspartic
66 proteinase activity ($p > 0.05$). The interaction between fluconazole, itraconazole, voriconazole,
67 posaconazole, isavuconazole and farnesol showed clear synergism (FICI ranges from 0.038 to
68 0.375) against one-day-old biofilms. Regarding *in vivo* experiments, daily 75 μM farnesol
69 treatment decreased the fungal burden in an immunocompromised murine model of
70 disseminated candidiasis, especially in case of inocula pre-exposed to farnesol ($p < 0.01$). In
71 summary, farnesol shows a promising therapeutic or adjuvant potential in traditional or
72 alternative therapies such as catheter lock therapy.

73
74 **Contribution to the field**

75 To date *Candida auris* had been reported from more than 35 countries on six different
76 continents. This newly emerged multidrug-resistant fungal pathogen causes nosocomial
77 outbreaks with high crude mortality rate. The number of resistant isolates to all of three main
78 classes of antifungals is steadily increasing worldwide. Due to the alarming emergence of
79 antifungal resistance, there is an urgent need to develop alternative antifungal therapies. An
80 attractive novel approach to combating fungal infections caused by multi-resistant pathogens
81 is treatments targeting quorum-sensing. Farnesol was the first described fungal quorum-
82 sensing molecule, causing hyphae-to-yeast transition in *C. albicans*. However, it is
83 noteworthy that farnesol may act differently in non-*albicans* *Candida* species; in addition, it
84 may have potent synergizing and/or antifungal effect. In this study, we performed
85 investigations focusing on growth, biofilm production ability, oxidative stress related enzyme
86 production, azole susceptibility and virulence in order to examine the effect of farnesol
87 exposure on *C. auris*. Based on our results, farnesol has an inhibitory effect both against
88 planktonic cells and biofilms. In addition, it showed a remarkable therapeutic potential in our
89 systemic immunocompromised mouse model. These results may support the development of
90 novel alternative therapies against *C. auris* infections in the future.

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1. Introduction

Candida auris is an emerging fungal pathogen causing outbreaks in healthcare settings with unacceptably high mortality rates ranging from 28% to 78% depending on the country (Jeffrey-Smith et al. 2018, Eyre et al. 2018). To date, 39 countries have reported *C. auris* associated infections (Jeffrey-Smith et al. 2018, Eyre et al. 2018, Kean et al. 2020). Based on last published data, the number of confirmed *C. auris* infections were 620 and 988 in Europe and United States of America, respectively (CDC 2019, ECDC 2018). Nosocomial *C. auris* outbreaks were reported from several countries including India, South Africa, Venezuela, Pakistan, and the United States (Vallabhaneni et al. 2016, Lockhart et al. 2017, Belkin et al. 2018). Previously, genetic analyses revealed more genetically unrelated clonal populations across three different continents. These clades are commonly classified as South African, South Asian, East Asian and South American clades (Lockhart et al. 2017). In addition, a recent study described a fifth *C. auris* clade in Iran from patient who never travelled outside that country (Abastabar et al. 2019, Chow et al. 2019).

Over 90% of clinical isolates are resistant to fluconazole whereas resistance to newer triazoles is variable (Dudiuk et al 2019, Romera et al 2019). The ratio of strains resistant to amphotericin B ranges from 8% to 50%, while echinocandin resistance remains infrequent (2% to 8%) (Dudiuk et al 2019). Alarmingly, isolates of *C. auris* with resistance to all three major antifungal classes have been reported in multiple countries including the USA (Ostrowsky et al. 2020). These multidrug-resistant strains may remain susceptible to nystatin and terbinafine (Sarma and Upadhyay 2017). *C. auris* biology have been extensively covered in recent papers (Rossato and Colombo 2018, Casadevall et al. 2019), however, the data about potential alternative treatment strategies remain scarce (Wall et al. 2019); therefore, there is an urgent need for the development of new antifungal therapies. In addition, multidrug-resistance is significantly more frequently reported in the case of *C. auris* biofilms (Kean and Ramage 2019). Thus, although the capacity to form biofilms is strain dependent in *C. auris*, they frequently pose a remarkable therapeutic challenge, especially because *C. auris* biofilms also have a considerable virulence capacity (Kean and Ramage 2019). Since data collected with *C. albicans* biofilms cannot be extrapolated to *C. auris* directly, such studies are urgently needed to meet this novel challenge (Kean and Ramage 2019).

Farnesol is a fungal quorum-sensing molecule that inhibits yeast-to-hyphae transition and promotes reverse morphogenesis in *C. albicans* (Hornby et al. 2001). Based on recent studies, farnesol acts synergistically with several antifungal agents against *C. albicans*, *C. glabrata*, *C. tropicalis* as well as against *C. parapsilosis* planktonic cells and/or biofilms (Katragkou et al. 2015, Kovács et al. 2016, Monteiro et al. 2017, Agustín et al. 2019), thus it has been proposed as a potential adjuvant therapeutic agent. In addition, its therapeutic potential has already been confirmed against *C. albicans* in murine models of mucosal infection (Hisajima et al. 2008, Bozó et al. 2016). Although farnesol is not beneficial in systemic infections caused by *C. albicans* (Navarhna et al. 2007), those data cannot necessarily be extrapolated to non-*albicans* species including *C. auris* (Semreen et al. 2019).

This study examines the effect of farnesol exposure on growth, biofilm production, oxidative stress-related enzyme production, triazole susceptibility and virulence of *C. auris*, in order to explore the background of the previously observed antifungal effect.

139 **2. Materials and methods**

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141 **2.1. Organisms**

142 Three *C. auris* isolates (isolates 10, 12, and 27) obtained from National Mycology Reference
143 Laboratory, United Kingdom were used together with the SC5314 *C. albicans* reference
144 strain. All three *C. auris* strains derived from the South Asian/Indian lineage (Borman et al.
145 2017). All *C. auris* isolates tested showed non-aggregating phenotype, which exhibit
146 comparable pathogenicity to that of *C. albicans* (Borman et al. 2016).

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148 **2.2. Toxicity experiments**

149 Ten μM , 50 μM , 150 μM and 300 μM farnesol were evaluated in terms of toxicity to the
150 Caco-2 cell line using a 3-(4,5-dimethyl-2-thiazolyl)-2,5-diphenyl-2H-tetrazolium bromide
151 (MTT) assay (Sigma, Budapest, Hungary) (Berridge et al. 2005). No toxicity was observed
152 with any concentration of farnesol.

153

154 **2.3. Growth related experiments for planktonic cells**

155 The effect of pre-exposure and continuous farnesol treatment on *C. auris* and *C. albicans*
156 planktonic cells was tested in RPMI-1640 (with L-glutamine and without bicarbonate, pH 7.0
157 with MOPS; Sigma, Budapest, Hungary) in two experimental settings: *i*) effect of various
158 farnesol concentrations against planktonic cells, *ii*) effect of various farnesol concentrations
159 against planktonic cells pre-exposed with farnesol (75 μM) for 24-hours. Seventy-five μM
160 farnesol was chosen as pre-exposure concentration because it corresponds to approximately
161 double the amount of physiological farnesol production of *C. albicans* (Weber et al. 2008).

162 Farnesol was obtained as 3M stock solution, which was diluted to a 30 mM working stock
163 solution in 100% methanol. The working concentrations of farnesol were prepared in RPMI-
164 1640 medium. Drug-free control was supplemented with 1% (vol/vol) methanol (Kovács et al.
165 2016, Bozó et al. 2016, Nagy et al. 2019). Farnesol concentrations tested were 10, 50, 100,
166 300 μM in all experiments.

167 Living cell number of planktonic cells was determined using time-kill experiments (Kovács et
168 al. 2014, Kovács et al. 2017). Briefly, samples (100 μL) were removed at 0, 2, 4, 6, 8, 10, 12
169 and 24 hours, serially diluted tenfold, plated (4 x 30 μL) onto Sabouraud dextrose agar and
170 incubated at 35 °C for 48 hours. All isolates were tested in three independent experiments and
171 the mean of the three values was used in the analysis. At given time points, one-way ANOVA
172 with Dunnett's post-testing was used to analyze the effect on living cell number exerted by
173 different farnesol concentrations compared to untreated control.

174

175 **2.4. Evaluation of extracellular phospholipase and aspartic proteinase activities exerted
176 by farnesol exposure**

177 Extracellular phospholipase production by farnesol-exposed (75 μM) and untreated *C. auris*
178 and *C. albicans* cells was examined on egg yolk medium (5.85% [wt/vol] NaCl, 0.05%
179 [wt/vol] CaCl₂, and 10% [vol/vol] sterile egg yolk [Sigma, Budapest, Hungary]). Aspartic
180 proteinase activity was evaluated on solid medium supplemented with bovine serum albumin
181 (0.02% [wt/vol] MgSO₄×7H₂O, 0.25% [wt/vol] K₂HPO₄, 0.5% [wt/vol] NaCl, 0,1% [wt/vol]
182 yeast extract, 2% [wt/vol] glucose and 0.25% [wt/vol] bovine serum albumin [Sigma,
183 Budapest, Hungary] agar medium). In case of both assay, 5 μL suspensions of 1×10^7
184 cells/mL were inoculated onto agar plates as described previously (Kantarcioglu and Yücel
185 2002). Colony diameters and precipitation zones (Pz) were measured after 7 days of
186 incubation at 35 °C (Price et al. 1982). Enzyme activities were measured in three independent
187 experiments for each isolate and are presented as means \pm standard deviations. Statistical
188 analysis of reactive species and enzyme production data were performed by paired Student's *t*

189 test using GraphPad Prism 6.05 software. The differences between values for treated and
190 control cells were considered significant if the p value was < 0.05 .

191

192 **2.5. Reactive species production and antioxidant enzyme activities exerted by farnesol** 193 **exposure**

194 Reactive species were measured in the presence or absence of one-day farnesol (75 μM)
195 exposure in RPMI-1640 by a reaction that converts 2',7'-dichlorofluorescein diacetate to 2',7'-
196 dichlorofluorescein (DCF) (Sigma, Budapest, Hungary) (Jakab et al. 2015, Jakab et al. 2019).
197 The amount of DCF produced is proportional to the quantity of reactive species. Catalase and
198 superoxide dismutase activities were determined as described previously by Jakab et al.
199 (2015) and Jakab et al. (2019). Reactive species and enzyme activities were measured in three
200 independent experiments for each isolate and are presented as means \pm standard deviations.
201 Statistical comparisons of reactive species and enzyme production data were performed by
202 paired Student's t test using GraphPad Prism 6.05 software. The differences between values
203 for treated and control cells were considered significant if the p value was < 0.05 .

204

205 **2.6. Susceptibility testing of planktonic cells to azoles and farnesol**

206 Antifungal susceptibility of *C. auris* isolates to fluconazole, itraconazole, voriconazole,
207 posaconazole, isavuconazole and to farnesol (all from Sigma, Budapest, Hungary) was tested
208 using the broth microdilution method in RPMI-1640 in line with the CLSI standard M27-A3
209 guideline (Clinical and Laboratory Standards Institute, 2008). The final concentrations of the
210 drug ranged between 0.5-32 mg/L, 0.008-0.5 mg/L and 1.17-300 μM mg/L for fluconazole,
211 other tested azoles and farnesol, respectively. Susceptibility testing for planktonic cells was
212 performed in 96-well microtitre plates at 35 $^{\circ}\text{C}$ for 24 hours. The inoculum was $0.5\text{-}2.5 \times 10^3$
213 cells/mL. Minimum inhibitory concentrations (MICs) were defined as at least 50% growth
214 reduction compared with untreated control. All isolates were tested in three independent
215 experiments and the median of the three values was used in the analysis.

216

217 **2.7. Biofilm formation**

218 *Candida* isolates were suspended in RPMI-1640 broth at a concentration of 1×10^6 cells/mL
219 and aliquots of 100 μl were inoculated onto flat-bottom 96-well sterile microtitre plates (TPP,
220 Trasadingen, Switzerland) and then incubated statically at 35 $^{\circ}\text{C}$ for 24 hours to produce one-
221 day-old biofilms (Pierce et al. 2008, Kovács et al. 2016).

222

223 **2.8. Metabolic activity changes of biofilms over time following farnesol exposure**

224 The effect of pre-exposure and continuous farnesol treatment on *C. auris* and *C. albicans*
225 biofilms was tested in three experimental settings: *i*) continuous farnesol treatment for 24-
226 hours during biofilm formation, *ii*) biofilm forming ability of cells pre-exposed with farnesol
227 (75 μM) for 24-hours prior to biofilm formation then continuously treated to given farnesol
228 concentrations for 24-hours during biofilm development, *iii*) effect of farnesol on one-day-old
229 biofilms. Farnesol concentrations tested were 10, 50, 100, 300 μM in all experiments.
230 Metabolic activity of sessile cells was determined at 0, 2, 4, 6, 8, 10, 12 and 24 hours using
231 XTT-reduction assay (Hawser 1996, Katragkou et al. 2015). All isolates were tested in three
232 independent experiments and the mean of the three values was used in the analysis. At given
233 time points, one-way ANOVA with Dunnett's post-testing was used to analyze the metabolic
234 activity change exerted by different farnesol concentrations compared to untreated control.
235 The differences between values for treated and control cells were considered significant if the
236 p value was lower than 0.05.

237

238 **2.9. Susceptibility testing of biofilms**

239 The activity of triazoles and farnesol against one-day-old biofilms was evaluated using the
240 XTT-assay (Hawser, 1996, Katragkou et al. 2014, Kovács et al. 2016, Nagy et al. 2019). The
241 concentrations tested in biofilm MIC determination ranged between 8-512 mg/L, 0.5-32
242 mg/L, 0.125-8 mg/L and 1.17-300 μ M for fluconazole, voriconazole/itraconazole,
243 posaconazole/isavuconazole and farnesol, respectively. To determine the 24-hour biofilm
244 MICs, one-day-old biofilms were first washed three times with 200 μ L sterile physiological
245 saline. All wells were filled with 100 μ L of 0.5 g/L XTT / 1 μ M menadione solution. The
246 plates were covered and incubated at 35 °C for 2 hours; afterwards, 80 μ L of the supernatant
247 was removed and transferred into a new sterile 96-well plate to measure the absorbance
248 spectrophotometrically at 492 nm. MICs were defined as the lowest concentration that
249 produced at least 50% reduction in metabolic activity of fungal biofilms compared to
250 untreated control (Katragkou et al. 2014, Kovács et al. 2016, Nagy et al. 2019). Three
251 independent experiments were performed for all isolates and the median of the three values
252 were presented.
253

254 **2.10. *In vitro* interactions between farnesol and azoles for planktonic cells and biofilms**

255 A fractional inhibitory concentration index (FICI) was used to evaluate drug-drug interactions
256 using a two-dimensional broth microdilution checkerboard assay both for planktonic and
257 sessile cells (Meletiadiis et al. 2005, Katragkou et al. 2015, Kovács et al. 2016). In the case of
258 *C. albicans*, combinations were tested only for biofilms because planktonic isolates are
259 generally susceptible to the tested azoles. The concentration ranges were as described above
260 for MIC determination against planktonic cells and biofilms. The FICI expressed as
261 Σ FIC= $FIC_A + FIC_B = MIC_A^{combination} / MIC_A^{alone} + MIC_B^{combination} / MIC_B^{alone}$, where MIC_A^{alone} and
262 MIC_B^{alone} are the MIC values of compounds A and B used alone and $MIC_A^{combination}$ and
263 $MIC_B^{combination}$ are the MICs of compounds A and B at the isoeffective combinations,
264 respectively. FICI was defined as the lowest Σ FIC (Meletiadiis et al. 2005, Katragkou et al.
265 2015, Kovács et al. 2016). The MIC values of the drugs alone and of all isoeffective
266 combinations were determined as the lowest drug concentrations showing at least 50%
267 reduction of turbidity for planktonic, or at least 50% reduction in metabolic activity of biofilm
268 compared to the untreated control cells. The interaction between azoles and farnesol was
269 interpreted as synergistic when FICI was ≤ 0.5 , as indifferent interaction when FICI was
270 between >0.5 and 4 and as antagonism when FICI was >4 (Meletiadiis et al. 2005, Katragkou
271 et al. 2015, Kovács et al. 2016).
272

273 **2.11. *In vivo* experiments**

274 BALB/c immunocompromised female mice (21-23 g) (Charles River) were used to examine
275 the effect of farnesol pre-exposure (75 μ M) and daily farnesol treatment (75 μ M) on virulence
276 of *C. auris* and compared to *C. albicans* SC5314. The animals were maintained in accordance
277 with the Guidelines for the Care and Use of Laboratory Animals. The experiments were
278 approved by the Animal Care Committee of the University of Debrecen, Debrecen, Hungary
279 (permission no. 12/2014 DEMÁB). Permanent immunosuppression was produced by
280 intraperitoneal administration of 150 mg/kg cyclophosphamide 4 days prior to infection, 100
281 mg/kg cyclophosphamide 1 day prior to infection, 100 mg/kg cyclophosphamide 2 days post-
282 infection and 100 mg/kg cyclophosphamide 5 days post-infection (Andes et al. 2010, Kovács
283 et al. 2014). In accordance with our preliminary experiments, mice were challenged
284 intravenously through the lateral tail vein; the infectious doses were 1×10^7 CFU/mouse and 8
285 $\times 10^3$ CFU/mouse in 0.2 mL volume for *C. auris* and *C. albicans*, respectively. Inoculum
286 density was confirmed by plating serial dilutions on Sabouraud dextrose agar (Kovács et al.
287 2014). Mice were divided into four groups (10 mice per group); *i*) untreated control mice; *ii*)
288 inoculation with 24 hours-long farnesol pre-exposed (75 μ M) cells; *iii*) there was no farnesol

289 pre-exposure to fungal cells prior to infection, but 75 μ M daily farnesol treatment
290 (corresponding to approximately 0.4 mg/kg) was started from 24 hours post-infection; *iv*) 24
291 hours-long farnesol pre-exposure (75 μ M) to fungal cells prior to infection; afterwards, 75 μ M
292 daily farnesol treatment was started at 24 hours post-infection.
293 Farnesol treatments were administered intraperitoneally in a volume of 0.5 mL. Control mice
294 were given 0.5 mL physiological saline intraperitoneally. At 6 days post-infection, mice were
295 euthanized, and their kidneys were removed (Fakhim et al. 2018), weighed and homogenized
296 aseptically. Fungal tissue burden was determined by quantitative culturing. Kidney tissue
297 burden was analyzed using Kruskal-Wallis test with Dunn's post-test (GraphPad Prism 6.05.).
298 Significance was defined as $p < 0.05$.

2.12. Histology

301 Kidneys of treated and untreated mice were subjected to histological investigations.
302 Histopathological examination and histochemical staining were performed on routine
303 formalin-fixed, paraffin-embedded mouse kidney tissues. Serial 4- μ m-thick sections were cut
304 from paraffin blocks, and Periodic acid-Schiff (PAS) staining was performed (Pupim et al.
305 2017, Kovács et al. 2019).
306

307 **3. Results**

308

309 **3.1. Effect of farnesol on *C. auris* and *C. albicans* planktonic cell growth**

310 Significant decrease was observed in growth rate of *C. auris* for 12 hours in the presence of
311 farnesol concentrations ranges from 50 to 300 μM both in case of farnesol unexposed and pre-
312 exposed cells ($p < 0.001-0.05$) (Figure 1A and B). At 24 hours, 100 and 300 μM farnesol
313 significantly decreased the viable cell count compared to untreated control in both
314 experimental settings ($p < 0.01-0.001$) (Figure 1A and B). Surprisingly, neither farnesol pre-
315 exposed nor unexposed *C. albicans* cells showed significant growth reduction at 24 hours
316 ($p > 0.05$) (Figure 1 C and D).

317

318 **3.2. Effects of farnesol on extracellular phospholipase and proteinase production of *C.*** 319 ***auris* and *C. albicans***

320 Farnesol treatment did not significantly influence the extracellular proteinase activity of either
321 *C. auris* or *C. albicans*. The Pz values were 0.83 ± 0.04 and 0.82 ± 0.05 for *C. auris* untreated
322 control and farnesol-exposed cells, respectively ($p > 0.05$), as compared to 0.53 ± 0.003 and
323 0.48 ± 0.02 with *C. albicans* untreated control and farnesol-exposed cells, respectively
324 ($p > 0.05$). Farnesol exposure resulted in significantly higher phospholipase activity for *C.*
325 *albicans* (Pz values were 0.48 ± 0.04 and 0.42 ± 0.02 for untreated control and farnesol-
326 exposed cells, respectively ($p < 0.01$); however, the Pz values were statistically comparable in
327 case of *C. auris* (Pz values were 0.9 ± 0.04 and 0.89 ± 0.05 for untreated control and farnesol-
328 exposed cells, respectively ($p > 0.05$)).

329

330 **3.3. Farnesol-induced oxidative stress and stress response in *C. auris* and *C. albicans*.**

331 Farnesol caused a significantly higher reactive species production in *C. auris* compared with
332 untreated control cells as presented in Table 1 ($p < 0.001$). This farnesol-related higher reactive
333 species level was associated with elevated superoxide dismutase ($p < 0.001$) but statistically
334 comparable catalase activity ($p > 0.05$) (Table 1). Farnesol treatment did not result in
335 significantly higher reactive species production in *C. albicans* ($p > 0.05$), which is in line with
336 the statistically comparable catalase and superoxide dismutase activity between farnesol
337 exposed cells and untreated control ($p > 0.05$) (Table 1).

338

339 **3.4. Effects of farnesol on biofilm forming ability and one-day-old biofilms of *C. auris*** 340 **and *C. albicans*.**

341 (i) *The effect of different farnesol concentrations on biofilm forming ability:* All tested
342 farnesol concentrations inhibited the metabolic activity of *C. auris* cells compared to control
343 cells at first 8 hours ($p < 0.001-0.05$); while, statistically comparable metabolic activities were
344 measured at 24 hours ($p > 0.05$) (Figure 2A). In contrast, all tested farnesol concentrations
345 inhibited the metabolic activity of *C. albicans* cells compared to untreated control at 24 hours
346 (Figure 2D).

347 (ii) *Biofilm forming ability of cells pre-exposed with farnesol for 24-hours (75 μM) prior to*
348 *biofilm formation:* Interestingly, we observed statistically significant differences in metabolic
349 activity of *C. auris* cells only at 24 hours between 50 and 300 μM (Figure 2B). In the case of
350 *C. albicans*, statistically significant differences in metabolic activity between 50 and 300 μM
351 were first observed at 8 hours (Figure 2E), but the metabolic activity of cells treated by
352 various concentrations was statistically comparable at 24 hours (Figure 2E).

353 (iii) *The effect of different farnesol concentrations against one-day-old biofilms:* Between 2
354 and 24 hours, 300 μM farnesol produced a potent anti-biofilm effect against *C. auris*
355 compared to control (Figure 2C). Interestingly, the low farnesol concentrations (10-50 μM)
356 increased the metabolic activity of *C. albicans* biofilms in the first 4 hours (Figure 2F).

357 However, the various farnesol treatments were statistically comparable against *C. albicans* at
358 24 hours (Figure 2F).

359

360 **3.5. Susceptibility results for planktonic cells and biofilms**

361 For *C. auris* isolates, the planktonic MICs ranged from 4 to >32 mg/L, from 0.03 to 0.06
362 mg/L, from 0.008 to 0.015 mg/L, from 0.015 to 0.03 mg/L and from 0.008 to 0.015 mg/L for
363 fluconazole, voriconazole, isavuconazole, itraconazole and posaconazole, respectively. The
364 susceptibility to fluconazole of isolate 10 was higher than the tentative fluconazole MIC
365 breakpoint (>32 mg/L) while the other two strains were susceptible to fluconazole (CDC
366 2020). In the case of planktonic *C. albicans* SC5314 reference strain, the median MIC values
367 were 0.125 mg/L, 0.015 mg/L, 0.015 mg/L, 0.125 mg/L and 0.008 mg/L for fluconazole,
368 voriconazole, isavuconazole, itraconazole and posaconazole, respectively. In case of biofilms,
369 the median MIC values are shown in Table 2.

370

371 **3.6. Interactions between triazoles and farnesol by FICI**

372 Only indifferent interactions were detected for planktonic cells of *C. auris* (data not shown).
373 The results of the triazole-farnesol interaction against one-day-old biofilms based on FICI are
374 summarized in Table 2. Antagonism was never observed. Synergy between triazoles and
375 farnesol was observed for all three *C. auris* isolates when grown in biofilm (FICI ranges from
376 0.038 to 0.375) (Table 2). For the *C. albicans* SC5314 strain, the interaction pattern observed
377 was very similar to *C. auris*; an indifferent interaction between an azole and farnesol was
378 observed only in case of fluconazole, although, the FICI value calculated was very close to
379 the synergy threshold (Table 2).

380

381 **3.7. *In vivo* experiments**

382 Results of the *in vivo* experiments are shown in Figure 3 and Figure 4 for *C. auris* and *C.*
383 *albicans*, respectively. Seventy-five μ M farnesol treatment decreased the fungal kidney
384 burden especially when farnesol pre-exposed *C. auris* cells were used as inoculum (Figure 3).
385 With *C. albicans*, all experimental settings resulted in statistically comparable kidney fungal
386 burdens compared to untreated control (Figure 4). The histopathology results observed were
387 in line with the fungal burden-related results. *C. auris* produced single yeast cells and
388 numerous budding yeast cells in untreated control mice. Although, inoculation by farnesol
389 pre-exposed cells caused large number of aggregates in kidney tissue; the daily farnesol
390 treatment markedly decreased the number of lesions (Figure 3). Both farnesol pre-exposure
391 and daily farnesol treatment caused several extended fungal lesions in kidney tissue in the
392 case of *C. albicans* infection (Figure 4), where single and budding yeast cells, pseudohyphae
393 and hyphae were observed in all groups (Figure 4).

394

395 4. Discussion

396 Only a few classes of antifungal agents are available for the treatment of fungal infections; in
397 addition, the antifungal drug discovery pipeline is slow and challenging, especially in case of
398 the newly emerging difficult-to-treat species such as *C. auris* (Roemer and Crysan 2014,
399 Scorzoni et al. 2017). Combination based therapeutic approaches have been proposed as
400 alternatives in recent years to treat the *C. auris* infections. The combination of flucytosine
401 with amphotericin B or micafungin may be relevant for the treatment of *C. auris* infections
402 (Bidaud et al. 2019). Moreover, synergistic interactions were observed between micafungin
403 and voriconazole (Fakhim et al. 2017).

404 The investigations of alternative/adjuvant treatments focusing on fungal quorum-sensing
405 molecules (e.g.: farnesol, tyrosol) have become an intensely researched area in recent years
406 (Mehmood et al. 2019). Several *in vitro* and *in vivo* studies were performed to evaluate the
407 antimicrobial effects of farnesol, which revealed that this compound may potentially serve as
408 an alternative or adjuvant drug (Jabra-Rizk et al. 2006, Hisajima et al. 2008, Katragkou et al.
409 2015, Kovács et al. 2016, Bozó et al. 2016, Nagy et al. 2019). Farnesol has a versatile effect
410 at physiological concentrations, however, the most prominent of these is its ability to
411 influence *C. albicans* morphology without markedly changing proliferation (Hornby 2001). It
412 is noteworthy that farnesol not only affects *C. albicans* but has a remarkable inhibitory effect
413 on other non-*albicans* species and moulds especially in suprphysiological concentrations
414 (Jabra-Rizk et al. 2006; Henriques et al. 2007; Rossignol et al. 2007; Weber et al. 2010,
415 Kovács et al. 2016). Our recent study reported that farnesol has a potential antifungal effect
416 against *C. auris* biofilms (Nagy et al. 2019), nevertheless, the physiological processes
417 underlying the observed antifungal activity of farnesol remain to be elucidated.

418 Farnesol did not affect the growth rate of planktonic *C. albicans*; but caused significant
419 reduction in growth rate in the case of *C. auris*. Moreover, farnesol inhibited the metabolic
420 activity of one-day-old biofilms in the first 24 hours, a phenomenon clearly absent with *C.*
421 *albicans*. The observed farnesol related effect in *C. albicans* is similar to those reported by
422 Hornby et al. (2001).

423 Farnesol has been suggested to modulate virulence, since it was shown to affect virulence-
424 associated phospholipase and aspartyl protease production in *C. albicans*. In this study,
425 farnesol exposure resulted in significantly higher phospholipase activity for *C. albicans*,
426 which is line with results reported by Fernandes et al. (2018). However, it did not enhance the
427 production of these enzymes in experiments with *C. auris*.

428 Farnesol was reported to cause a dose-dependent production of reactive species and could
429 increase resistance to oxidative stress in *C. albicans* (Davis-Hanna et al. 2008, Deveau et al.
430 2010, Giacometti et al. 2011), which is concordant with our results. However, farnesol
431 treatment resulted in a significant increase of reactive species production in *C. auris*, resulting
432 in an elevated level of superoxide dismutase but not catalase, demonstrating that farnesol
433 might not contribute to protection against oxidative stress in *C. auris*. Such stress-related
434 differences between *C. albicans* and *C. auris* were also observed previously with other
435 stressor compounds. *C. auris* was more resistant to hydrogen-peroxide compared to *C.*
436 *albicans*; but it was less tolerant to the superoxide-generating agent menadione and the tert-
437 butyl hydroperoxide, and moreover displayed significantly higher resistance to cationic stress
438 imposed by either sodium chloride or calcium chloride compared to *C. albicans* (Day et al.
439 2018).

440 To date, catheter-associated infections caused by *C. auris* have been reported by several
441 authors, which are attributable to the previously well-documented biofilm-forming ability of
442 this species (Dewaele et al. 2018). Previous studies reported the frequency of central line
443 infections by *C. auris* to be between 11% and 92%. (Taori et al 2019, Schelenz 2016, Lee
444 2011). Although sessile communities show significantly higher resistance to the majority of

445 frequently used antifungals compared to planktonic susceptibilities (Kean and Ramage 2019),
446 the efficacy of such antifungal agents can be enhanced using adjuvants such as farnesol (Nagy
447 et al. 2019). A clear synergy between the tested triazoles and farnesol against *C. auris*
448 biofilms was demonstrated, similarly to the combinations of echinocandins and farnesol
449 (Nagy et al. 2019). Farnesol modulates the expression of genes linked to ergosterol
450 biosynthesis, which may explain the synergy of this compound with triazoles (Yu et al. 2012).
451 Although the *in vitro* effect of farnesol is well known especially against *C. albicans*, its *in*
452 *vivo* role remains controversial and raises several questions. Navarathna et al. (2007) showed
453 that exogenous farnesol (20 mM/mouse) can enhance the pathogenicity of *C. albicans*,
454 increasing the mortality in a murine model of systemic candidiasis. In contrast, Hisajima et al.
455 (2008) observed a farnesol-induced protective effect (at a dose 9 μ M/mouse) in *C. albicans*-
456 associated oropharyngeal candidiasis. Although Boz o et al. (2016) revealed that farnesol
457 alone is not protective in a murine vulvovaginitis model (150-300 μ M/mouse), it did enhance
458 the fluconazole activity against a fluconazole-resistant *C. albicans* isolate. In addition,
459 chitosan nanoparticles containing miconazole and farnesol also inhibited fungal proliferation
460 in a mouse vulvovaginitis model at ≥ 240 μ M (Fernandes et al. 2019). To the best of our
461 knowledge, there is no reported data concerning the *in vivo* activity of farnesol against non-
462 *albicans Candida* species. In this study, daily farnesol treatment decreased the *C. auris* fungal
463 burden in mouse kidneys regardless of previous farnesol exposure of the inoculum. In
464 addition, in the case of inocula pre-exposed to farnesol, the reduction of fungal cell numbers
465 was statistically significant, which is concordant with our *in vitro* growth-related results. The
466 antifungal activity observed may be explained by the elevated levels of reactive species
467 previously measured *in vitro*, which could not be detected in equivalent experiments with *C.*
468 *albicans*. Furthermore, the amphiphilic properties of farnesol allows for its integration into
469 cell membranes, affecting membrane fluidity and integrity (Bringmann et al. 2000; Funari et
470 al. 2005; Jabra-Rizk et al. 2006; Scheper et al. 2008). Farnesol was shown to affect cellular
471 polarization and membrane permeability in *C. parapsilosis* and *C. dubliniensis* (Jabra-Rizk et
472 al. 2006; Rossignol et al. 2007), which may also explain the observed antifungal effect in our
473 study. However, it is noteworthy that the inoculation of farnesol pre-exposed cells without
474 daily farnesol treatment resulted in a more virulent *C. auris* population and increased fungal
475 burden. The 24-hours-long pre-exposure without further continuous treatment of farnesol may
476 influence the expression of virulence determinants or membrane properties similar to
477 fluconazole pre-treatment, which may explain the virulence enhancer effect reported
478 previously (Navarathna et al. 2005).
479 In conclusion, our results clearly demonstrate farnesol-related differences in physiology
480 between *C. albicans* and *C. auris*. Based on our *in vivo* studies, farnesol has a remarkable
481 therapeutic potential against *C. auris*; in addition, it reverses the well-documented resistance
482 to newer triazoles reported for *C. auris* biofilms. However, further genome-wide gene
483 expression analysis with *C. auris* is needed in order that each aspect of farnesol-related effects
484 (e.g.: short-term exposure vs. long-term exposure) can be elucidated.

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492

493

494 **6. Declaration of interest**

495

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500 **7. References**

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789 **Table 1** Farnesol-induced oxidative stress response in *C. auris* and *C. albicans*
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Oxidative stress related parameter	Untreated cultures		Farnesol-exposed cultures	
	<i>C. auris</i>	<i>C. albicans</i>	<i>C. auris</i>	<i>C. albicans</i>
Catalase [kat (kg protein) ⁻¹]	1.41 ± 0.03	0.60±0.07	1.56±0.09	0.48±0.07
SOD [munit (mg protein) ⁻¹]	85.69±5.42	78.13±4.51	170.11±17.37***	81.41±6.12
DCF [nmol DCF (OD ₆₄₀) ⁻¹]	3.96±0.89	9.69±1.01	23.54±4.51***	11.45±1.15

792
 793 Mean ± standard deviation values calculated from three independent experiments are
 794 presented.

795
 796 *** Significant differences at $p < 0.001$, as calculated by the paired Student's *t*-test compared to
 797 untreated control and farnesol-treated cultures for *C. auris*.

798 **Table 2** Minimum inhibitory concentration of fluconazole (FLU), voriconazole (VOR),
799 itraconazole (ITRA), posaconazole (POSA) and isavuconazole (ISA) alone and in
800 combination with farnesol (FAR) against *C. auris* (10, 12 and 27) and *C. albicans* SC5314
801 biofilms (sMIC). Furthermore, *in vitro* interactions by fractional inhibitory concentration
802 index (FICI) determination of fluconazole, voriconazole, itraconazole, posaconazole and
803 isavuconazole in combination with farnesol against *C. auris* and *C. albicans* biofilms. Median
804 MIC values and FICI values from three independent experiments are presented.

Isolates	Median sMIC values				Interaction analysis	
	sMIC alone		sMIC in combination		Median FICI	Type of interaction
	FLU (mg/L)	FAR (µM)	FLU (mg/L)	FAR (µM)		
10	>512 ^a	300	64	75	0.375	Synergy
12	>512 ^a	300	64	75	0.35	Synergy
27	>512 ^a	300	64	75	0.375	Synergy
SC5314	>512 ^a	150	64	75	0.56	Indifferent
	VOR (mg/L)	FAR (µM)	VOR (mg/L)	FAR (µM)		
10	64	150	0.5	4.69	0.093	Synergy
12	64	300	0.5	4.69	0.061	Synergy
27	64	300	0.5	9.38	0.038	Synergy
SC5314	16	150	1	4.69	0.09	Synergy
	ITRA (mg/L)	FAR (µM)	ITRA (mg/L)	FAR (µM)		
10	16	300	0.5	4.69	0.155	Synergy
12	32	300	0.5	9.375	0.140	Synergy
27	16	300	0.5	9.375	0.123	Synergy
SC5314	8	150	0.5	4.69	0.187	Synergy
	POSA (mg/L)	FAR (µM)	POSA (mg/L)	FAR (µM)		
10	16	150	0.25	2.34	0.062	Synergy
12	16	150	0.25	2.34	0.062	Synergy
27	16	150	0.25	2.34	0.062	Synergy
SC5314	2	150	0.25	4.69	0.28	Synergy
	ISA (mg/L)	FAR (µM)	ISA (mg/L)	FAR (µM)		
10	4	300	0.125	9.38	0.091	Synergy
12	8	300	0.125	18.75	0.062	Synergy
27	4	300	0.125	9.38	0.091	Synergy
SC5314	8	150	0.5	4.69	0.28	Synergy

839 ^a MIC is off-scale at >512 mg/l, 1024 mg/l (one dilution higher than the highest tested
840 concentration) was used for analysis

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843

844 **Figure 1**

845 Time-kill curves of farnesol against *C. auris* (A and B) and *C. albicans* (C and D) isolates in
846 RPMI-1640 for farnesol unexposed (A and C) and farnesol pre-exposed (B and D) cells (75
847 μM), respectively. Each timepoint represents mean \pm SEM (standard error of mean) of cell
848 count derived from isolates.

849

850 **Figure 2**

851 Metabolic activity changes over time in case of biofilm formation in the presence of given
852 farnesol concentrations (10-300 μM) for *C. auris* (A) and *C. albicans* (D), respectively.
853 Metabolic activity changes over time in case of biofilm formation by farnesol pre-exposed
854 cells (75 μM) in the presence of given farnesol concentrations (10-300 μM) for *C. auris* (B)
855 and *C. albicans* (E), respectively. Metabolic activity changes over time for one-day-old
856 preformed biofilms in the presence of given farnesol concentrations (10-300 μM) for *C. auris*
857 (C) and *C. albicans* (F), respectively. Each time-point represents mean \pm SEM (standard error
858 of mean) of metabolic activity of clinical isolates (three independent experiments per isolate).

859

860 **Figure 3**

861 The kidney burden of *C. auris* in a systemically infected mouse model. The bars represent the
862 means \pm SEM (standard error of mean) of kidney tissue burdens of BALB/c mice. Significant
863 differences between CFU numbers were determined based on comparison with the untreated
864 controls. Levels of significant differences are indicated (** $p < 0.01$). Histological changes in
865 kidney tissue from mice suffering from systemic candidiasis with or without farnesol
866 treatment in the presence or absence of farnesol pre-exposure were examined by Periodic
867 acid-Schiff staining.

868

869 **Figure 4**

870 The kidney burden of *C. albicans* in a systemically infected mouse model. The bars represent
871 the means \pm SEM (standard error of mean) of kidney tissue burdens of BALB/c mice.
872 Significant differences between CFU numbers were determined based on comparison with the
873 untreated controls. Histological changes in kidney tissue from mice suffering from systemic
874 candidiasis with or without farnesol treatment in the presence or absence of farnesol pre-
875 exposure were examined by Periodic acid-Schiff staining.