

Candida albicans biofilms are generally devoid of persister cells

Iryna Denega, Christophe d'Enfert, Sophie Bachellier-Bassi

▶ To cite this version:

Iryna Denega, Christophe d'Enfert, Sophie Bachellier-Bassi. Candida albicans biofilms are generally devoid of persister cells. Antimicrobial Agents and Chemotherapy, American Society for Microbiology, 2019, 63 (5), pp.1-9. 10.1128/AAC.01979-18. pasteur-02086290

HAL Id: pasteur-02086290 https://hal-pasteur.archives-ouvertes.fr/pasteur-02086290

Submitted on 1 Apr 2019

HAL is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers. L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.



Distributed under a Creative Commons Attribution - NonCommercial - ShareAlike | 4.0 International License

- 1 *Candida albicans* biofilms are generally devoid of persister cells
- 2
- 3 Iryna Denega^{1,2}, Christophe d'Enfert¹ and Sophie Bachellier-Bassi^{1,#}
- 4
- ⁵ ¹Institut Pasteur, INRA, Unité Biologie et Pathogénicité Fongiques, 25 rue du Docteur
- 6 Roux, Paris, France
- 7 ²Univ. Paris Diderot, Sorbonne Paris Cité, Cellule Pasteur, Paris, France
- 8
- 9
- ¹⁰ [#]Corresponding author: Sophie Bachellier-Bassi, Institut Pasteur, Unité Biologie et
- 11 Pathogénicité Fongiques, Département Mycologie, 25 rue du Docteur Roux, F-75015
- 12 Paris, France; Phone: +33 (1) 45 68 86 19; E-mail: sophie.bachellier-
- 13 <u>bassi@pasteur.fr</u>
- 14
- 15
- 16
- 17 RUNNING TITLE: Lack of persister cells in C. albicans biofilms
- 18 KEYWORDS: Candida albicans, biofilms, antifungal tolerance, persistence
- 19

20 ABSTRACT

21 Candida albicans is known for its ability to form biofilms - communities of 22 microorganisms embedded in an extracellular matrix developing on different 23 surfaces. Biofilms are highly tolerant to antifungal therapy. This phenomenon has 24 been partially explained by the appearance of so-called persister cells, phenotypic 25 variants of wild-type cells, capable of surviving very high concentrations of 26 antimicrobial agents. Persister cells in C. albicans were found exceptionally in 27 biofilms while none were detected in planktonic cultures of this fungus. Yet, this topic 28 remains controversial as others could not observe persister cells in biofilms formed 29 by the *C. albicans* SC5314 laboratory strain. Due to ambiguous data in the literature, 30 this work aimed to re-evaluate the presence of persister cells in *C. albicans* biofilms. We demonstrated that isolation of C. albicans "persister cells" as described 31 32 previously was likely to be the result of survival of biofilm cells that were not reached 33 by the antifungal. We tested biofilms of SC5314 and its derivatives, as well as 95 34 clinical isolates, using an improved protocol, demonstrating that persister cells are 35 not a characteristic trait of C. albicans biofilms. Although some clinical isolates are 36 able to yield survivors upon the antifungal treatment of biofilms, this phenomenon is 37 rather stochastic and inconsistent.

38

40 **INTRODUCTION**

The yeast *Candida albicans* is a commensal of humans but also one of the most prevalent fungal pathogens, responsible for superficial infections as well as lifethreatening systemic infections (1). *C. albicans* is recognized for its ability to form biofilms that are most frequently associated with nosocomial infections, particularly in immunocompromised patients.

46 C. albicans biofilms are communities of microorganisms with a complex structure 47 composed of different cell types embedded in an extracellular matrix (2-4). They develop on different types of surfaces, either living or inert, and are characterized by 48 49 their high tolerance to antifungals. The latter can result from the properties of the 50 extracellular matrix that can serve as a trap for drug molecules (5-7). An additional 51 source of antifungal tolerance has been proposed to result from the occurrence in 52 biofilms of so-called persister cells, a subpopulation of phenotypic variants of wild-53 type cells, capable of surviving concentrations of antimicrobial agents well above the Minimal Inhibitory Concentration (MIC) (8). Persister cells were first described in 54 55 bacterial cultures as a drug tolerant subpopulation, that upon removal of the 56 antimicrobial agent gave rise to a new population of susceptible cells (9). Persisters 57 are known to be genetically identical to the rest of the population, thus persistence is 58 a non-inherited trait (10–12).

In the clinical setting, persisters are usually associated with relapse of infections and with the development of chronic infections. For bacterial persisters, several mechanisms and pathways involved in their development have been described (13).

In 2006, LaFleur et al. have presented the first report of persister cells in biofilms of *C. albicans*, which could contribute to biofilm tolerance to antifungals (8). In their
paper, the authors have reported that *C. albicans* exhibit a biphasic killing curve,

65 when exposed to the antifungals such as amphotericin B (AMB), chlorhexidine or the 66 combination of both. This phenomenon is explained by the presence of a multidrugtolerant subpopulation of persister cells within a biofilm, while planktonic cultures of 67 68 C. albicans were found to be devoid of persisters. Notably, the experiments for this 69 study were performed using in vitro biofilm model of C. albicans, developed in 70 polystyrene 96-well plates. Following this work and relying on the protocol for 71 persister cells isolation described therein (8), persister cells in C. albicans biofilms 72 were described by a few other groups (14–16). However, later work by the Douglas 73 group showed that not all Candida species and strains were able to form persister 74 cells in laboratory-grown biofilms (17). This was in particular the case for C. albicans 75 strain SC5314 (18), the parental strain of almost all C. albicans strains used for 76 functional genomics and molecular genetics studies. Unlike in the previously 77 mentioned papers (8, 14-16), the protocol Al-Dhaheri and Douglas (17) used for 78 persisters isolation involved growing biofilms on silicone discs followed by their immersion into an antifungal solution. As the topic of C. albicans persister cells 79 80 remains controversial, the main objective of this work was to re-evaluate their 81 occurrence in *in vitro*-grown *C. albicans* biofilms.

82

83 METHODS

84 Strains and growth conditions

In this study we used 3 reference strains (listed in Table 1) and a set of 95 C.

- 86 albicans, 3 C. tropicalis and 3 C. parapsilosis clinical isolates (Table S1).
- Yeast precultures were grown overnight in YPD (1% yeast extract, 2% peptone, 2%
 glucose) with shaking at 30°C.

Biofilms were grown either in RPMI 1640 medium with L-glutamine (buffered with 50 mM HEPES), as described in (8) and (20), or in GHAUM medium (SD supplemented with 2% glucose and 1 mg/mL histidine, 1 mg/mL arginine, 0.02 mg/mL uridine and 2 mg/mL methionine (21)).

Resistance was checked on solid YNG (6,7 g/L yeast nitrogen base without amino
acids and with ammonium sulfate, 2% glucose and 2% agar) supplemented with 10
µg/mL AMB.

96

97 Biofilm growth and persister cells isolation

98 To assess persister cell appearance in biofilms we used two protocols adapted either 99 from (8) or (14). The first protocol uses 96-well plates and the biofilms are grown in 100 RPMI. In the second protocol the biofilms are grown in 24-well plates but using 101 GHAUM medium instead of YNB.

102 Biofilm growth

103 Overnight cultures were washed in sterile 1x PBS and diluted in the corresponding 104 medium to $OD_{600} 0.3$. Either 100 µL or 1 mL of cells in the 96-well plate or the 24-well 105 plate, respectively, were allowed to adhere for 1.5 h without agitation. The non-106 adhered cells were then washed with 1X PBS, the same volume of fresh medium 107 was added, plates were covered with a breathable seal and biofilms were allowed to 108 form for 48 h at 37°C with agitation (110 rpm) with a medium change after 24 hours.

109 Antifungal treatment

110 Media were carefully aspirated from the 48 h-old biofilms, without disrupting the 111 biofilm structure. Biofilms were washed once with either 100 μ L or 1 mL of 1x PBS, 112 respectively, and treated with a 100 μ g/mL AMB solution in either RPMI or GHAUM 113 for 24 hours at 37°C, statically. AMB solutions were prepared from an 8 mg/mL stock

in DMSO, so that the final concentration of DMSO in a working solution did not
exceed 1.25%. For control biofilms, corresponding amount of DMSO was added to
the medium instead of the antifungal solution.

117 This step was either performed using the same volumes of antifungal solution as for 118 biofilm growth as described in (8) and (14) or increasing the volume of antifungal to

119 fill the well up to the top (350 µL or 3 mL for 96- and 24-well plates, respectively).

120 Clinical isolates were first treated with 64 μ g/mL AMB solution. Strains giving rise to 121 colonies were then tested 5 times with 100 μ g/mL AMB.

122 Plating

Upon 24 hours of antifungal treatment, AMB solution was aspirated and biofilms were washed twice with 1X PBS prior to plating on YPD-agar plates. Biofilms were resuspended in 1x PBS/0.05% Tween-20. For the AMB-treated samples, the whole biofilms were plated. For control biofilms, serial dilutions were performed to allow CFU counting. CFU were counted after incubating the plates at 30°C for 48 h.

128

129 **RESULTS AND DISCUSSION**

130 In this work, we aimed to study the occurrence of persister cells in *C. albicans* 131 biofilms. We applied the protocol published by LaFleur and colleagues, growing the 132 biofilms in RPMI and in a 96-well plate format (8). We set up the protocol with 3 133 *C. albicans* prototroph strains, namely SC5314, CEC369 and CEC4664 - prototroph 134 derivatives of BWP17 (22) and SN76 (23), respectively. BWP17 and SN76 are 135 independent auxotroph derivatives of SC5314.

We encountered a technical problem at the biofilm recovery step, usually performed by scraping the cells in 1x PBS and vortexing prior to plating (8, 14, 16, 24). In our hands, the cells could not be properly resuspended and plated, as clumps of the

biofilms would usually remain stranded inside the tips. Consequently, the CFU
numbers obtained were highly variable for all samples, making any further analysis
and comparison impossible (data not shown).

142 We decided to test alternative approaches to circumvent the stickiness of biofilms. 143 Resuspending cells in 20% glycerol/1X PBS for plating helped reducing stickiness, 144 but did not improve consistency (data not shown). We hypothesized that EDTA might 145 reduce adherence of biofilms by binding bivalent cations that are required for the 146 activity of cell surface adhesins (25). Thus, we attempted applying 20% glycerol with 147 a range of EDTA concentrations (0, 50, 100 mM) for plating. 100 µL of EDTA 148 solutions of different concentrations were added to biofilms and left for 10 minutes at 149 room temperature prior to biofilm disruption by scraping and vortexing. None of the 150 applied EDTA solutions allowed abolishing stickiness. Additionally, colonies growing 151 on YPD-agar exhibited a wrinkled morphology, most probably linked to the toxicity of 152 EDTA (26). Finally, we tried adding Tween-20 (0.05%) to PBS. Tween-20 eradicated the problems of stickiness and poor disruption and improved recovery of cells from 153 154 the biofilms (Fig. 1). The effect on cell viability was tested using a planktonic culture 155 of SC5314 that was washed and plated on YPD-agar using PBS and PBS-Tween-20 156 solutions. No impact on viability was observed (data not shown). Thus, in the 157 experiments described below, biofilms were resuspended in a 0.05% Tween-20/1X 158 PBS solution.

However, even after this modification, the ratio of cells that survived AMB treatment was still inconsistent between repeats. According to Lafleur and colleagues the ratios of *C. albicans* persister cells in biofilms vary from 0.1% to 2% for different strains, notably from 0.05 to 0.1% for strain CAI4 – a derivative of *C. albicans* SC5314 (8). Our values hardly ever exceeded 0.01% persisters per biofilm, even after improving

the recovery protocol, thus bordering with statistical error. We reasoned that increasing the surface of a biofilm and changing the growth media could improve persister yields and decided to test the protocol described in (14), applying the modifications that were mentioned previously. However, the problem of inconsistency and low ratios of persisters remained (Fig. 2).

169 In all protocols described previously, the volumes of the media and solutions used for 170 biofilm growth, washing, and AMB treatment were identical. Upon a careful 171 observation, we noticed that C. albicans cells form a dense rim at the border of the 172 air and liquid phases, as a result of agitation during growth. Treating a biofilm with the 173 exact same volume of antifungal and growth medium in static conditions thus could 174 result in cells from the rim escaping treatment. We decided to increase the volume of 175 the applied antifungal solution (filling wells to the top) and, to our surprise, this 176 change in the protocol led to a complete eradication of persisters for the laboratory 177 strain SC5314 and its derivatives. Reproducibly, we did not get any persisters after 178 applying this change for all strains for both RPMI- and GHAUM-grown biofilms. Thus, 179 the volume of the antifungal applied in the original protocols for persister isolation 180 was skewing the results. Increasing the volume of antifungal eliminated this bias, 181 resulting in a complete eradication of any survivors after the antifungal treatment.

In our work we used a modified protocol for persister cells isolation with a starting cell suspension of OD_{600} 0.3 used for biofilm growth instead of 0.1 as described in the original protocols (8, 14). To assess the impact of the initial cell number used for seeding biofilms on persister cells' appearance, we tested our protocol for SC5314 using cell suspensions of OD_{600} 0.1, 0.3 and 0.5 for seeding. Regardless of the initial biomass, persister cells did not form in SC5314 biofilms grown either in RPMI or GHAUM (data not shown).

189 These results made us question the very existence of persister cells in C. albicans 190 biofilms. Previously, Al-Dhaheri and Douglas showed that not all strains of 191 C. albicans can form persister cells (17). Particularly, in their hands, SC5314 biofilms 192 lost all viability after exposure to 30 µg/mL AMB. However, biofilms of another clinical 193 isolate, GDH2346, appeared to contain a small proportion (0.01%) of cells that 194 survived 100 µg/mL AMB treatment. These authors used a different in vitro model for 195 assessing persistence, as they grew biofilms on silicone disks that were transferred 196 to a new well filled with an antifungal solution. This prevented an escape of any cells 197 from the antifungal treatment. Thus, our modified protocol for treatment of biofilms 198 formed in 96-well or 24-well plates corroborated the results obtained by the Douglas 199 group for C. albicans strain SC5314 (17).

200 Since the clinical isolate GDH2346 could give rise to survivors (17), we could not 201 exclude that persisters could emerge in biofilms of different C. albicans isolates. 202 Additionally in 2010, LaFleur and colleagues isolated and described C. albicans 203 strains from patients with long-term oral infection, that gave yield to increased levels 204 of persisters (up to 8.9%) (24). These were called hip-mutants, by analogy with the 205 high persister strains previously described for bacteria (27, 28). Although hip-mutants 206 were identified using a protocol that showed limitations in our hands, we 207 hypothesized that some C. albicans clinical isolates could generally be more prone to 208 form persisters than others (namely SC5314). To test this assumption, we tested 95 209 clinical isolates (Table S1) for their ability to form biofilms and the occurrence of 210 persister cells following AMB treatment. In a first round of experiments, biofilms were 211 treated with a 64 µg/mL AMB solution. Only 38 isolates (39.6%) displayed survivors 212 (notably, never exceeding a rate of 0.02%). According to the generally accepted 213 concept of persistence (10), the frequency of persisters' appearance is independent

214 of the increase in antibiotic concentration. Thus in a second round of experiments, 215 biofilms were developed for these 38 isolates and treated with a 100 µg/mL AMB 216 solution. Notably, only 7 isolates out of these 38 displayed survivors when grown with 217 100 µg/mL AMB (CEC3622, CEC3668, CEC3669, CEC4514, CEC4521, CEC5317, 218 CEC5318). These 7 strains, together with 4 other isolates randomly picked in the 219 remaining 31 strains (CEC712, CEC3708, CEC3711, CEC5316), were tested seven 220 more times with 100 µg/mL of AMB. In most cases these strains did not yield 221 persister cells (Fig. 3); however, 7 strains (CEC3622, CEC3669, CEC4514, 222 CEC4521, CEC5316, CEC5317 and CEC5318) gave rise to small numbers of 223 survivors in one to four of the experiments (Fig. 3), with the survival rate never 224 exceeding 9.1*10⁻⁴% per biofilm (for CEC3622). This could be explained either by the 225 stochastic nature of persistence as a phenomenon or by technical errors during the 226 experiment.

We tested up to 30 randomly picked colonies for three isolates (CEC3622, CEC4514 and CEC5316) on YNG medium containing 10 µg/mL of AMB. None of the tested colonies was able to grow in presence of amphotericin B (data not shown), proving that their survival was not a result of AMB resistance development.

231 With an improved protocol in our hands, we decided to test other Candida species for 232 their ability to form persister cells in biofilms. Previously, Al Dhaheri and Douglas (17) 233 reported that clinical isolates of C. krusei (Glasgow strain) and C. parapsilosis (AAHB 234 4479) developed persister cells in biofilms (approximately 0.001% and 0.07%, 235 respectively) upon treatment with 100 µg/mL AMB. We selected 3 clinical isolates of 236 C. tropicalis (CEC5296, CEC5297, CEC5298) and 3 of C. parapsilosis (CEC5299, 237 CEC5300, CEC5301) from our lab collection to test with our protocol. One of the C. 238 tropicalis strains (CEC5298), as well as the 3 selected C. parapsilosis strains were

unable to grow as biofilms, and were excluded from the study. *C. tropicalis* CEC5296 and CEC5297 formed proper biofilms, with a small fraction of persisters varying between $2*10^{-5}$ - $6.4*10^{-3}$ % and $2.3*10^{-7}$ - $2.6*10^{-4}$ % respectively (data not shown). Such low values are comparable to the survival rates we observed for some of the *C. albicans* clinical isolates tested in this study. As before, we cannot exclude that these survivors are persister cells arising within *C. tropicalis* biofilms, or that they are the consequence of a technical error during the experiment.

246

247 CONCLUSION

248 Since 1944, when Bigger first described persister cells in Staphylococcus (9), many 249 advances have been made in exploring this phenomenon, especially in bacteria. It is 250 known that microbial cultures growing in vivo can sometimes be very difficult to 251 eradicate completely by an antibiotic treatment, causing relapses or development of 252 chronic infections in patients. From an evolutionary point of view, a small pool of cells 253 with the same genotype as the rest of the population but differing in their ability to 254 tolerate stress - including drug treatment - provides a form of insurance to the 255 population.

The phenomenon of persistence has not only been described for bacteria, but also in other types of pathogens, and it has been proposed that persister cells significantly contributed to the recalcitrance of *C. albicans* biofilms to antifungal treatments (29– 31).

C. albicans persister cells were first described in 2006 (8), and since then just a handful of reports, sometimes contradictory, have been presented. In our study, we explored standard protocols to obtain persisters, and showed that their proportion in biofilms formed by different *C. albicans* strains has been overestimated. Only Al-

264 Dhaheri and Douglas did not detect persisters in SC5314 biofilms (17). In their study, 265 biofilms were grown on silicon discs that were transferred in antifungal solutions for 266 treatment. In contrast, the other published experiments were performed using 96-well 267 plates and RPMI medium, or 24-well plates and SD-based medium, while keeping 268 the incubation volumes constant throughout the experiment (8, 14–16). In this study, 269 we modified the latter protocols (8, 14-16) by increasing the volume of antifungal. 270 This change led to the eradication of biofilms, indicating that previously detected 271 "persisters" were likely the result of survival of cells that were not reached by the 272 antifungal. Our results corroborate the findings of Al-Dhaheri and Douglas (17).

Notably, these authors were able to detect some persisters in biofilms of a clinical isolate (17), but the ratio obtained was much lower (0.01%) than the numbers published by others (8, 14). Although some of the clinical isolates of *C. albicans* and *C. tropicalis* tested in our study were occasionally able to yield survivors after the treatment of biofilms with AMB, this phenomenon was rather inconsistent, pointing either to the stochastic nature of persistence itself, or another skew in the protocol while carrying out particular experiments.

At this time, we cannot completely exclude the possibility of persistence in all *C. albicans* strains, though with the described protocol we managed to disprove their presence for 92 *C. albicans* strains out of 98. It is important to stress that our results reflect only the behaviour of *C. albicans* biofilms grown *in vitro*; we cannot rule out that in the context of the host, persister cells could appear and contribute to the general resistance and dissemination of *C. albicans*.

286

287 ACKNOWLEDGEMENTS

Iryna Denega is part of the Pasteur - Paris University (PPU) International PhD Program. This project has received funding from the European Union's Horizon 2020 research and innovation programme under the Marie Sklodowska-Curie grant agreement N° 665807, and from the Institut Carnot Pasteur Microbes & Santé. This work has been supported by grants from the French Government's Investissement d'Avenir program (Laboratoire d'Excellence Integrative Biology of Emerging Infectious Diseases, ANR-10-LABX-62-IBEID) to C.d'E.

295

296 **REFERENCES**

Sardi JCO, Scorzoni L, Bernardi T, Fusco-Almeida AM, Mendes Giannini MJS.
 2013. *Candida* species: current epidemiology, pathogenicity, biofilm formation,
 natural antifungal products and new therapeutic options. J Med Microbiol 62:10–24.

Ramage G, Saville SP, Thomas DP, López-Ribot JL. 2005. *Candida biofilms*:
an update. Eukaryot Cell 4:633–638.

302 3. Nobile CJ, Johnson AD. 2015. *Candida albicans* biofilms and human disease.
303 Annu Rev Microbiol 69:71–92.

Chandra J, Kuhn DM, Mukherjee PK, Hoyer LL, McCormick T, Ghannoum
 MA. 2001. Biofilm formation by the fungal pathogen *Candida albicans*: development,
 architecture, and drug resistance. J Bacteriol 183:5385–5394.

5. Nett JE, Crawford K, Marchillo K, Andes DR. 2010. Role of Fks1p and matrix glucan in *Candida albicans* biofilm resistance to an echinocandin, pyrimidine, and polyene. Antimicrob Agents Chemother 54:3505–3508.

6. Nett JE, Sanchez H, Cain MT, Andes DR. 2010. Genetic basis of *Candida*biofilm resistance due to drug sequestering matrix glucan. J Infect Dis 202:171–175.

7. Nett JE, Sanchez H, Cain MT, Ross KM, Andes DR. 2011. Interface of *Candida albicans* biofilm matrix-associated drug resistance and cell wall integrity
regulation. Eukaryot Cell 10:1660–1669.

8. LaFleur M. 2006. *Candida albicans* bofilms produce antifungal-tolerant
 persister cells. Antimicrob Agents Chemother 50:3839–3846.

317 9. Bigger J. 1944. Treatment of staphylococcal infections with penicillin by
318 intermittent sterilisation. The Lancet 244:497–500.

319 10. Brauner A, Fridman O, Gefen O, Balaban NQ. 2016. Distinguishing between
320 resistance, tolerance and persistence to antibiotic treatment. Nat Rev Microbiol
321 14:320–330.

11. Lewis K. 2007. Persister cells, dormancy and infectious disease. Nat Rev
Microbiol 5:48–56.

12. Lewis K. 2010. Persister cells. Annu Rev Microbiol 64:357–372.

13. Harms A, Maisonneuve E, Gerdes K. 2016. Mechanisms of bacterial
 persistence during stress and antibiotic exposure. Science 354:aaf4268.

14. Li P, Seneviratne CJ, Alpi E, Vizcaino JA, Jin L. 2015. Delicate metabolic
 control and coordinated stress response critically determine antifungal tolerance of
 Candida albicans biofilm persisters. Antimicrob Agents Chemother 59:6101–6112.

15. Truong T, Zeng G, Qingsong L, Kwang LT, Tong C, Chan FY, Wang Y,
Seneviratne CJ. 2016. Comparative ploidy proteomics of *Candida albicans* biofilms
unraveled the role of the *AHP1* gene in the biofilm persistence against amphotericin
B. Mol Cell Proteomics 15:3488–3500.

16. Sun J, Li Z, Chu H, Guo J, Jiang G, Qi Q. 2016. *Candida albicans*amphotericin B-tolerant persister formation is cosely related to surface adhesion.
Mycopathologia 181:41–49.

17. Al-Dhaheri RS, Douglas LJ. 2008. Absence of amphotericin B-tolerant
 persister cells in biofilms of some *Candida species*. Antimicrob Agents Chemother
 52:1884–1887.

340 18. Gillum AM, Tsay EY, Kirsch DR. 1984. Isolation of the *Candida albicans* gene
341 for orotidine-5'-phosphate decarboxylase by complementation of *S. cerevisiae ura3*342 and *E. coli pyrF* mutations. Mol Gen Genet 198:179–182.

19. Enjalbert B, Rachini A, Vediyappan G, Pietrella D, Spaccapelo R, Vecchiarelli
A, Brown AJP, d'Enfert C. 2009. A Multifunctional, synthetic *Gaussia princeps*luciferase reporter for live imaging of *Candida albicans* infections. Infect Immun
77:4847–4858.

20. De Brucker K, De Cremer K, Cammue BPA, Thevissen K. 2016. Protocol for
determination of the persister subpopulation in *Candida albicans* biofilms. Methods
Mol Biol 1333:67–72.

21. Cabral V, Znaidi S, Walker LA, Martin-Yken H, Dague E, Legrand M, Lee K,
Chauvel M, Firon A, Rossignol T, Richard ML, Munro CA, Bachellier-Bassi S,
d'Enfert C. 2014. Targeted changes of the cell wall proteome influence *Candida albicans* ability to form single- and multi-strain biofilms. PLoS Pathog 10:e1004542.

Wilson RB, Davis D, Mitchell AP. 1999. Rapid hypothesis testing with *Candida albicans* through gene disruption with short homology regions. J Bacteriol 181:1868–
 1874.

357 23. Noble SM, Johnson AD. 2005. Strains and strategies for large-scale gene
 358 deletion studies of the diploid human fungal pathogen *Candida albicans*. Eukaryot
 359 Cell 4:298–309.

24. LaFleur MD, Qi Q, Lewis K. 2010. Patients with long-term oral carriage harbor
 high-persister mutants of *Candida albicans*. Antimicrob Agents Chemother 54:39–44.

362 25. Klotz SA, Rutten MJ, Smith RL, Babcock SR, Cunningham MD. 1993.
 363 Adherence of *Candida albicans* to immobilized extracellular matrix proteins is
 364 mediated by calcium-dependent surface glycoproteins. Microb Pathog 14:133–147.

26. Chudzik B, Malm A, Rautar B, Polz-Dacewicz M. 2007. *In vitro* inhibitory
activity of EDTA against planktonic and adherent cells of *Candida* sp. Ann Microbiol
57:115.

27. Moyed HS, Bertrand KP. 1983. *hipA*, a newly recognized gene of *Escherichia coli* K-12 that affects frequency of persistence after inhibition of murein synthesis. J
Bacteriol 155:768–775.

28. Wolfson JS, Hooper DC, McHugh GL, Bozza MA, Swartz MN. 1990. Mutants
of *Escherichia coli* K-12 exhibiting reduced killing by both quinolone and beta-lactam
antimicrobial agents. Antimicrob Agents Chemother 34:1938–1943.

374 29. Borghi E, Borgo F, Morace G. 2016. Fungal Biofilms: update on resistance, p.
375 37–47. In Fungal Biofilms and related infections. Springer, Cham.

376 30. Ramage G, Rajendran R, Sherry L, Williams C. 2012. Fungal biofilm
377 resistance. Int J Microbiol 2012: 1-14.

378 31. Mathé L, Dijck PV. 2013. Recent insights into *Candida albicans* biofilm
379 resistance mechanisms. Curr Genet 59:251–264.

32. Ropars J, Ropars J, Maufrais C, Diogo D, Marcet-Houben M, Perin A, Sertour
N, Mosca K, Permal E, Laval G, Bouchier C, Ma L, Schwartz K, Voelz K, May RC,
Poulain J, Battail C, Wincker P, Borman AM, Chowdhary A, Fan S, Kim SH, Pape
PL, Romeo O, Shin JH, Gabaldon T, Sherlock G, Bougnoux M-E, d'Enfert C. 2018.
Gene flow contributes to diversification of the major fungal pathogen *Candida albicans*. Nature Communications 9:2253.

38. 33. Bougnoux M-E, Morand S, d'Enfert C. 2002. Usefulness of multilocus
sequence typing for characterization of clinical isolates of *Candida albicans*. J Clin
Microbiol 40:1290–1297.

389 34. Odds FC, Bougnoux M-E, Shaw DJ, Bain JM, Davidson AD, Diogo D,
390 Jacobsen MD, Lecomte M, Li S-Y, Tavanti A, Maiden MCJ, Gow NAR, d'Enfert C.
391 2007. Molecular Phylogenetics of *Candida albicans*. Eukaryot Cell 6:1041–1052.

392 35. Bougnoux M-E, Kac G, Aegerter P, d'Enfert C, Fagon J-Y, CandiRea Study
393 Group. 2008. Candidemia and candiduria in critically ill patients admitted to intensive
394 care units in France: incidence, molecular diversity, management and outcome.
395 Intensive Care Med 34:292–299.

396 36. Sdoudi K, Bougnoux M-E, Hamoumi RE, Diogo D, Mdaghri NE, d'Enfert C,
397 RazkiA. 2014. Phylogeny and Diversity of *Candida albicans* Vaginal Isolates from
398 Three Continents. Int J Curr Microbiol App Sci. 3:471–480.

399 37. Schönherr FA, Sparber F, Kirchner FR, Guiducci E, Trautwein-Weidner K, 400 Gladiator A, Sertour N, Hetzel U, Le GTT, Pavelka N, d'Enfert C, Bougnoux M-E, 401 Corti CF, LeibundGut-Landmann S. 2017. The intraspecies diversity of *C. albicans* 402 triggers qualitatively and temporally distinct host responses that determine the 403 balance between commensalism and pathogenicity. Mucosal Immunol 10:1335– 404 1350.

38. Shin JH, Bougnoux M-E, d'Enfert C, Kim SH, Moon C-J, Joo MY, Lee K, Kim
M-N, Lee HS, Shin MG, Suh SP, Ryang DW. 2011. Genetic Diversity among Korean *Candida albicans* Bloodstream Isolates: Assessment by Multilocus Sequence Typing
and Restriction Endonuclease Analysis of Genomic DNA by Use of BssHII. J Clin
Microbiol 49:2572–2577.

39. Bougnoux M-E, Diogo D, François N, Sendid B, Veirmeire S, Colombel JF,
Bouchier C, Van Kruiningen H, d'Enfert C, Poulain D. 2006. Multilocus sequence
typing reveals intrafamilial transmission and microevolutions of *Candida albicans*isolates from the human digestive tract. J Clin Microbiol 44:1810–1820.

40. Bougnoux M-E, Aanensen DM, Morand S, Théraud M, Spratt BG, d'Enfert C.
2004. Multilocus sequence typing of *Candida albicans*: strategies, data exchange
and applications. Infect Genet Evol 4:243–252.

417 41. Dieng Y, Sow D, Ndiaye M, Guichet E, Faye B, Tine R, Lo A, Sylla K, Ndiaye
418 M, Abiola A, Dieng T, Ndiaye JL, Le Pape P, Gaye O. 2012. Identification of three
419 *Candida africana* strains in Senegal. J Mycol Med 22:335–340

42. Garnaud C, Botterel F, Sertour N, Bougnoux M-E, Dannaoui E, Larrat S,
421 Hennequin C, Guinea J, Cornet M, Maubon D. 2015. Next-generation sequencing
422 offers new insights into the resistance of *Candida* spp. to echinocandins and azoles.
423 J Antimicrob Chemother 70:2556–2565.

424

425

426

427 FIGURE LEGENDS

428

Fig. 1 Effect of Tween 20 on the recovery of CFUs from *C. albicans* SC5314 biofilms. *C. albicans* SC5314 was allowed to form biofilms in 100 μL RPMI in a 96-well plate according to the protocol adapted from (8). Error bars: standard deviation (SD) of 6 biological replicates generated from 2 independent experiments. # - non-significant difference, *** - significant difference, p=0,0007 (unpaired t-test was applied to compare datasets).

435

436	Fig. 2. Schemes of the protocols (A) and levels of persisters (B) obtained
437	from biofilms grown using modified protocol from (14). Biofilms were grown in 1
438	mL of GHAUM medium in 24-well plates before application of either 1 mL of AMB
439	solution (on the left) or 3 mL of AMB solution (on the right). Ratios of surviving cells
440	are as follow: SC5314 - 5.6*10 ⁻⁴ %, CEC369 - 2.6*10 ⁻⁵ %, CEC4664 - 9.4*10 ⁻⁵ %.
441	Error bars: SD of 6 biological replicates generated from 2 independent experiments.
442	
443	
444	Fig. 3. Analysis of persister cell formation in 11 clinical isolates. Biofilms were
445	grown in 1 mL of GHAUM medium in 24-well plates, and treated with 3 mL of AMB

solution (modified protocol from (14)). The values obtained from 7 biofilms were used

to draw the graph.

448

449 TABLE 1. *C. albicans* reference strains used in this study

STRAIN	GENOTYPE	REFERENCE
SC5314		(18)
CEC369	ura3::λimm434/ura3::λimm434 ARG4/arg4::hisG HIS1/his1∆::hisG RPS1/RPS1::Clp10	(19)
CEC4664	ura3∆::λimm434/ura3∆::λimm434 iro1∆::λimm434/iro1∆::λimm434 ADH1/adh1::P _{TDH3} - carTA::SAT1 arg4Δ/ARG4 his1∆::hisG/HIS1 RPS1/RPS1::Clp10	Lab's collection

TABLE 2. Clinical isolates used in this study

NAME	REFERENCE	
CEC704	(32)	
CEC712	(32)	
CEC718	(32)	
CEC723	(32)	
CEC1289	(33)	
CEC1424	(34)	
CEC2018	(35)	
CEC2019	(34)	
CEC2020	(36)	
CEC2021	(35)	
CEC2022	(37)	
CEC2871	(38)	
CEC2876	(38)	
CEC3494	(39)	
CEC3533	(37)	
CEC3534	(36)	
CEC3535	(36)	
CEC3536	(36)	
CEC3540	(32)	
CEC3541	(33)	
CEC3544	(39)	
CEC3547	(39)	
CEC3548	(39)	
CEC3549	(39)	
CEC3550	(40)	
CEC3553	(39)	
CEC3555	(33)	
CEC3556	(36)	
CEC3560	(34)	
CEC3561	(39)	
CEC3596	(33)	
CEC3611	(36)	
CEC3614	(36)	
CEC3615	(36)	
CEC3621	(32)	
CEC3622	(36)	
CEC3623	(32)	

CEC3626	(39)
CEC3627	(36)
CEC3634	(36)
CEC3637	(34)
CEC3659	(35)
CEC3662	(35)
CEC3663	(32)
CEC3664	(35)
	(35)
CEC2669	(35)
	(35)
	(35)
CEC3672	(35)
CEC3675	(35)
CEC3681	(35)
CEC3682	(35)
CEC3685	(35)
CEC3706	(35)
CEC3708	(35)
CEC3711	(35)
CEC4035	(32)
CEC4039	(32)
CEC4103	(41)
CEC4104	(41)
CEC4106	(41)
CEC4108	(41)
CEC4256	(42)
CEC4259	(42)
CEC4481	(32)
CEC4482	(32)
	(32)
CEC4403	(32)
CEC4400	(32)
	(32)
	(32)
CEC4489	(32)
CEC4492	(32)
CEC4494	(32)
CEC4495	Lab's collection
CEC4496	(32)
CEC4501	Lab's collection
CEC4504	Lab's collection
CEC4505	Lab's collection
CEC4511	Lab's collection
CEC4514	Lab's collection
CEC4515	Lab's collection
CEC4517	Lab's collection
CEC4521	Lab's collection
CEC4524	Lab's collection
CEC4526	(32)
CEC4527	Lab's collection
CEC4547	Lab's collection
CEC4548	Lab's collection
CEC4549	Lab's collection
CEC4550	Lah's collection
CEC4552	(32)
0004002	(J∠) (22)
0100029	(52)

CEC5316	Lab's collection
CEC5317	Lab's collection
CEC5318	Lab's collection





