

1 **New generation modified azole antifungals against multidrug-resistant *Candida auris***

2 Yiyuan Chen¹, Yunxiao Li¹, Charlotte K. Hind², Md. Mahbub Hasan^{1, 4}, Kazi S. Nahar^{1,3},
3 Melanie Clifford², J. Mark Sutton^{1, 2 *}, Khondaker Miraz Rahman^{1 *}

4

5 ¹Institute of Pharmaceutical Science, King's College London, 150 Stamford Street, London.
6 SE1 9NH, United Kingdom.

7 ²UK Health Security Agency, Vaccine Evaluation and Development Centre, Manor Farm Road,
8 Salisbury. SP4 0JG, United Kingdom.

9 ³Department of Natural Sciences, University of Middlesex, The Burroughs, Hendon, London
10 NW4 4BT

11 ⁴Department of Genetic Engineering and Biotechnology, Faculty of Biological Sciences,
12 University of Chittagong, Chattogram, 4331, Bangladesh

13

14 **Abstract**

15 The increasing prevalence of antifungal resistance and the limited numbers of antifungal
16 agents available to treat patients with invasive fungal disease underscores the urgent need for
17 novel drug classes. *Candida auris* has emerged as a major public pathogen of global concern
18 with reduced effective treatment options. A targeted modification of azole core-scaffold with
19 a cyclic heteroaliphatic linker linked aromatic and heteroaromatic rings resulted in compounds
20 with exceptional activity, with MICs ranging from 0.016 to 4 µg/mL against a panel of *Candida*
21 *auris*, including azole-resistant isolates. The research employed a systematic approach,
22 varying substitutions of the linkers and the terminal aromatic rings to develop a structure
23 activity relationship (SAR). The compounds also showed excellent activity against public health
24 *Candida* species including *C. albicans*, *N. glabrata*, *C. tropicalis* and *C. parapsilosis*, with MICs
25 less than 1 µg/mL for most compounds. The study identified compounds **7**, **18** and **21** as
26 promising lead candidates with superior potency than both fluconazole and voriconazole
27 against both *C. auris* and diverse *Candida* strains. The compounds were found to be non-toxic
28 up to 50 mg/Kg in a *Galleria mellonella* model while showing efficacy against *C. auris* at a dose
29 of 5 mg/Kg. This study offers a new chemical scaffold that can be taken forward to develop
30 new generation azole antifungal agents against *C. auris*.

31

32 Introduction

33 Antifungal resistance is the ability of a fungus to grow and survive in the presence of antifungal
34 drugs. This can lead to severe infections that are hard to treat.^{1, 2} Antifungal drugs are used to
35 treat a variety of fungal infections, including candidiasis, aspergillosis, and cryptococcosis.³⁻⁵
36 Resistance can occur naturally or develop over time due to exposure to antifungal drugs or
37 fungicides. Improper use of antifungal drugs, such as low doses or short courses, can also
38 contribute to resistance.^{2, 6, 7} Some fungi, like *Aspergillus* and certain *Candida* species, are
39 resistant to some or all types of antifungal drugs.⁸⁻¹¹ Among them, *Candida auris* is a new and
40 highly resistant fungus that can spread quickly in healthcare settings.¹²⁻¹⁴

41 *C. auris* is an emerging fungal pathogen that is resistant to many antifungal drugs. It can cause
42 serious infections in hospitals and other healthcare settings.¹⁵⁻¹⁸ The US Centers for Disease
43 Control and Prevention (CDC) and the World Health Organization (WHO) both consider *C. auris*
44 to be a major threat to public health.¹⁹ It was first discovered in Japan in 2009 and has since
45 been reported in over 47 countries worldwide, with 6 clades emerging.²⁰ A study of patients
46 with echinocandin-susceptible *C. auris* bloodstream infection at three hospitals in Brooklyn,
47 New York, found that 30.1% of patients died within 30 days and 44.6% died within 90 days.²¹

48 Resistant *Candida* strains are relatively prevalent in clinical settings. Collected data from the
49 CDC showed that around 7% of the clinical *Candida* strains isolated from hospitalised patients
50 suffering from bloodstream infections exhibited resistance to marketed antifungal drugs.^{22, 23}
51 In another study approximately 90% of the *C. auris* isolates were found to be resistant to at
52 least one commercially available antifungal drug, and 30% of the clinical strains were non-
53 susceptible to more than one antifungal on the market.²⁴ All the data from public health
54 authorities makes it clear that innovation in the treatment of antifungal infections, particularly
55 the drug resistant *C. auris* is urgently necessary.

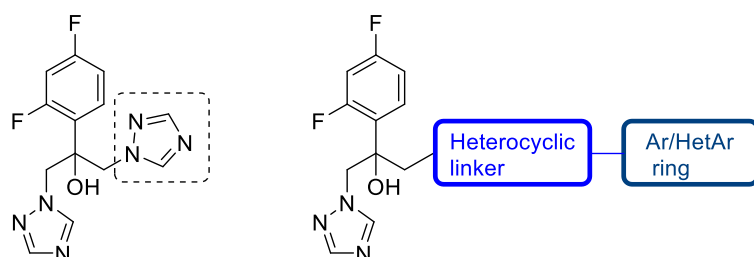
56 There are only four major classes of antifungal drugs available on the market for treating
57 systemic fungal infections, namely azoles (e.g., fluconazole, voriconazole, itraconazole, and
58 posaconazole), polyenes (typically amphotericin B), echinocandins (e.g. micafungin and
59 caspofungin), and pyrimidine analogues (mainly flucytosine or its salt form).^{25, 26} Azoles
60 (imidazoles and triazoles, containing two and three nitrogens on theazole ring, respectively)
61 are a class of antifungal drugs widely used to treat fungal infections in humans.²⁷ They work
62 by inhibiting the synthesis of ergosterol, an essential component of fungal cell membranes.
63 This disruption weakens the cell membrane and leads to the death of the fungal pathogen.^{28,}
64 ²⁹

65 Despite the potential of azoles, the increasing levels of resistance against them is becoming a
66 concern in tackling the healthcare burden caused by fungal infections. The treatment of fungal
67 pathogens using azoles triggers the upregulation of efflux pumps in *Candida spp.*, including
68 CDR1, MDR1, RDC3, SNQ2, and YHD3 in *C. auris*.³⁰⁻³² Moreover, reports show that *Candida*
69 *spp.* can also overexpress azole targets such as the ERG11 gene (that encodes lanosterol 14-
70 alpha-demethylase),^{33, 34} sequestrations of azoles within vacuole and biofilms (in fluconazole-
71 resistant *C. albicans*),^{35, 36} and by modification of the targets by mutation over time (ERG11
72 mutation triggered by the application of azoles in *Candida spp.*).^{37, 38} To develop azoles that

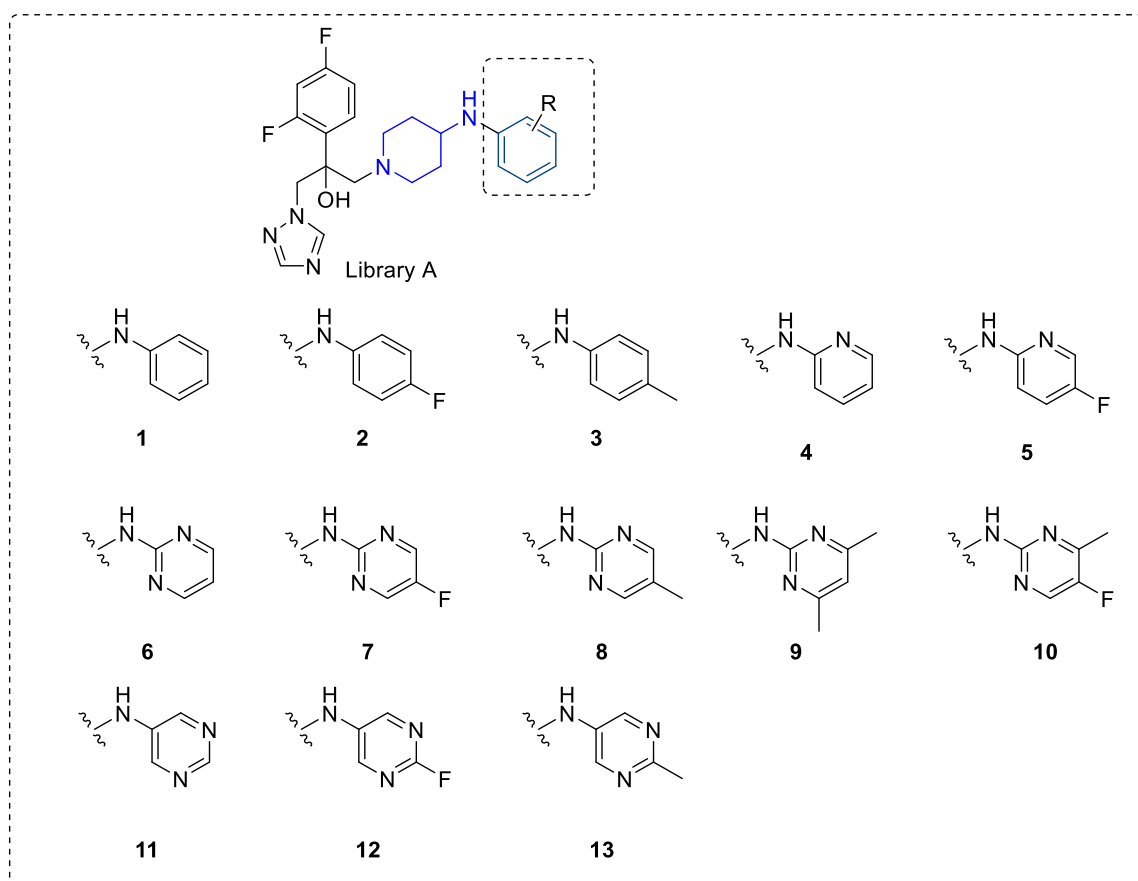
73 can overcome resistance observed in *C. auris* and other drug-resistant *Candida* species, we
 74 made a targeted modification in the azole core scaffold with various types of hetero-aliphatic
 75 linkers with different lengths, sizes, and ring strains (Figure 1A), and modified the terminal
 76 aromatic and heteroaromatic ring. This medicinal chemistry led approach has led to
 77 compounds with activity against multidrug-resistant *C. auris* and other clinically important
 78 *Candida* species that can be taken forward towards pre-clinical development.

79

(A)



(B)



80

81 **Figure 1.** (A) general structure of the modified azole compounds. (B) compounds of library A
 82 containing a piperidine linker and different aromatic or heteroaromatic rings.

83

84

85 Results and Discussion

86 Design of modified azole compounds

87 We aimed to design fluconazole analogues with improved interaction with the fungal
88 lanosterol 14 α -demethylase (LDM) which is the target for azole antifungals and is involved in
89 the biosynthesis of ergosterol.³⁹ For all compounds, one of the triazole rings of fluconazole
90 was replaced by a heteroaliphatic linker connected with an aromatic or heteroaromatic ring
91 (Figure 1A). For library A compounds, we designed compounds with a six-membered
92 heterocyclic piperidine ring as the linker as it has been previously been used to generate azole
93 modifications and compounds with this linker have shown MICs comparable to fluconazole^{40,}
94 ⁴¹. The piperidine ring was substituted at the 4-position with an amino group which was
95 considered as a point of diversity, and was substituted with a number of six-membered
96 aromatic and heteroaromatic rings to establish the structure activity relationship (Figure 1B).

97

98 **Table 1.** The comparative binding affinity of novel azole analogues and commercially available
99 azoles with fungal target lanosterol 14 α -demethylase (LDM).

Compound	<i>C. albicans</i>		<i>C. auris</i>	
	Chem Score	ΔG (kcal/mol)	Chem Score	ΔG (kcal/mol)
Fluconazole	18.16	-18.94	16.07	-18.8
Voriconazole	24.6	-26.72	21.03	-26.48
1	40.46	-42.6	36.16	-39.65
2	41.22	-42.69	37.32	-38.34
3	36.3	-37.52	32.25	-33.7
4	34.39	-37.4	33.68	-34.13
5	31.77	-33.34	30.77	-33.33
6	32.88	-35.63	30.96	-32.1
7	41.22	-42.69	37.32	-38.34
8	39.45	-41.6	35.6	-34.54
9	40.8	-37.6	37.6	-37.4
10	38.6	-36.54	38.45	-38.33
11	34.31	-35.65	30.44	-33.61
12	33.45	-33.66	31.32	-34.7
13	35.74	-37.4	33.22	-35.6

100

101 Designed compounds having phenyl, pyridine and pyrimidine as terminal aromatic or
102 heteroaromatic fragments were then utilised for the computational analysis against the target
103 LDM from different *Candida* species along with the clinically relevant drugs Fluconazole and

104 Voriconazole (Table 1). The data from the computational analysis showed that the newly
105 designed modified fluconazole compounds showed better binding affinity to LDM enzyme
106 from both *C. albicans* and *C. auris* compared to both fluconazole and voriconazole. In terms
107 of both ChemScore and Gibbs free energy ΔG most of the modified azole compounds showed
108 almost twice the binding affinity compared to Fluconazole. The results of binding affinity
109 remained similar across both species of *Candida* (Table 1).

110

111

112 **Synthesis of the library A**

113 This library consisted of 13 compounds in a two-step process, the fragments were synthesised
114 or purchased (compounds **2b**, **4b** – **7b**, **11b**) and then connected to the azole core by an
115 epoxide ring-opening reaction to form the final compounds. Fragments were synthesised by
116 following three different approaches based on the electronic environment of the connecting
117 carbon (Scheme 1).

118 The fragments were synthesised either by reductive amination (**1a**, **2a**, **12a** and **13a**) or by
119 simple S_NAr reactions depending on the electronic environment of the terminal
120 aromatic/heteroaromatic ring (**Scheme 1A**). In the case of fragment **10a**, a palladium-
121 catalysed organometallic reaction was carried out to obtain the fragment. The epoxide core
122 of the azole compounds was synthesised using the classic Corey-Chaykovsky reaction
123 conditions (Scheme 1B). The final compounds were synthesised by reacting the amine
124 fragments **1b-13b** with the epoxide core 14 with a nucleophilic epoxide ring opening reaction.
125 This resulted in racemic final compounds **1** to **13** which were evaluated for their antifungal
126 activity (Scheme 1).^{42, 43}

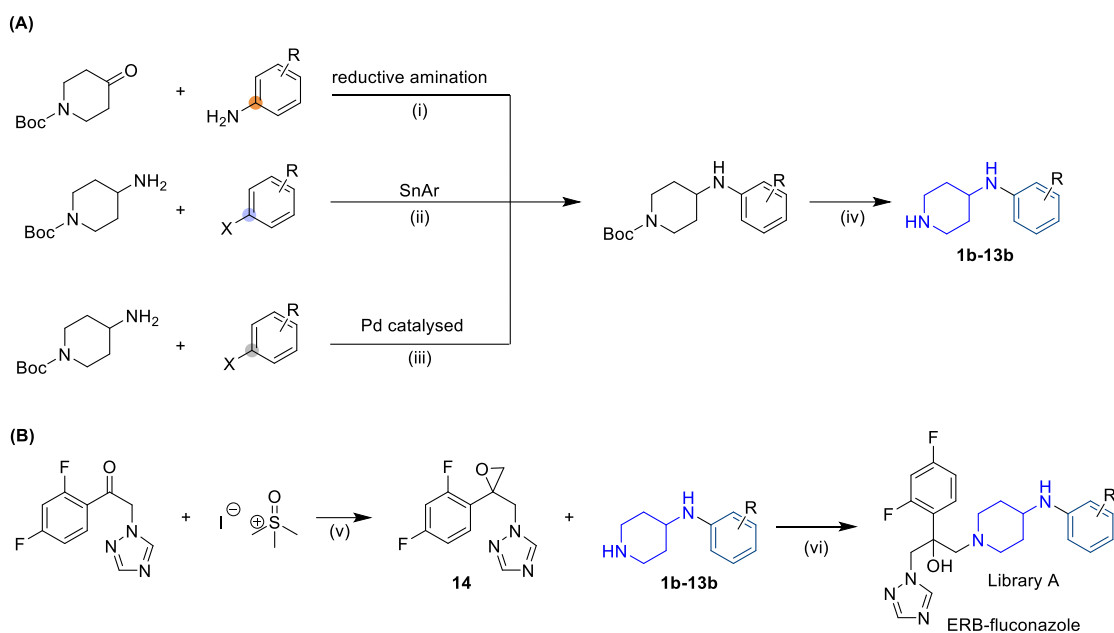
127

128

129

130

131

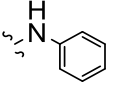
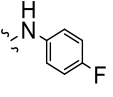
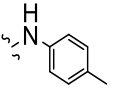
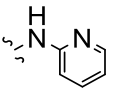
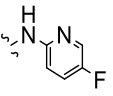
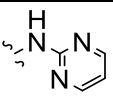


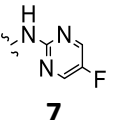
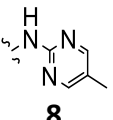
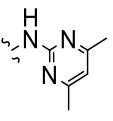
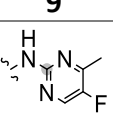
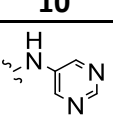
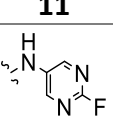
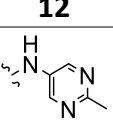
132

133

134 **Scheme 1.** General synthetic routes for Library A where different aromatic/heteroaromatic
 135 fragments (A) were connected to the fluconazole core via a piperidine linker (B). Conditions:
 136 (i) NaBH(OAc)₃, AcOH, DCM / MeOH, overnight; (ii) DIPEA, MeCN, DMA, 160 – 180 °C, 2 – 3 h
 137 or TEA, ethylene glycol, 200 °C, 15 min; (iii) rac-BINAP, Pd₂(dba)₃, KOTu, toluene, 100 °C,
 138 overnight; (iv) 4M HCl in 4-dioxane, r.t., 1 – 2 h; (v) NaOH (aq.), Toluene, 80 °C, Microwave 50
 139 min; (vi) TEA, EtOH, 80 °C, overnight.

140 **Table 2.** Antifungal activity ($\mu\text{g/mL}$) of modified fluconazole compounds against *Candida*, and their corresponding LogP value.

	Log P	<i>C. albicans</i>	<i>C. glabrata</i>	<i>C. tropicalis</i>	<i>C. parapsilosis</i>	<i>C. auris</i>							
		NCPF3281	NCPF8018	NCPF8760	NCPF3209	TDG1912	TDG2512	TDG1102	TDG2211	TDG2506	NCPF8984	NCPF8971	NCPF8977
Fluconazole	0.87	0.06 – 0.125	4	16	0.25	> 128	8	64	64	128	128	32	16
 1	2.49	0.12	0.12	0.12 – 0.5	0.12	0.12 – 0.25	0.03	0.5	0.25	1	1	0.0625	0.0625
 2	2.65	0.12	0.12	1	0.12	0.5 – 1	0.06	0.5	0.01563	2	4	0.0625	0.0625
 3	2.98	0.002 – 0.004	0.125	0.0625	< 0.001	0.002 – 0.016	0.0156	0.0625	1	0.0625	0.125	< 0.002 – 0.5	< 0.001
 4	1.87	≤ 0.25	4	2	≤ 0.25	2	0.125	2	0.25	4	4	0.125	0.25
 5	2.03	≤ 0.25	0.5 – 2	0.25 – 0.5	≤ 0.25	≤ 0.25	0.06	0.5	0.5	2	2	0.0625	0.25
 6	1.06	0.12	0.12 – 0.25	0.5	0.12	1	0.125	2	1	4	4	0.25	0.5

Log P	Chemical Structure	<i>C. albicans</i>	<i>C. glabrata</i>	<i>C. tropicalis</i>	<i>C. parapsilosis</i>	<i>C. auris</i>							
		NCPF3281	NCPF80 18	NCPF876 0	NCPF3209	TDG1 912	TDG2 512	TDG1 102	TDG2 211	TDG2 506	NCPF89 84	NCPF89 71	NCPF89 77
1.22		≤ 0.125	0.25	0.25	≤ 0.125	0.25	0.06	4	0.5	≤ 0.03	≤ 0.03	0.25	≤ 0.03
1.55		≤ 0.125	0.25	1	≤ 0.125	0.5	0.25	1	16	8	8	0.5	0.25
2.47		≤ 0.06 – 0.125	1	1	≤ 0.06	0.25 – 1	0.0625	0.5	0.25	0.25	2	0.0625	0.25
1.93		≤ 0.125	0.25	0.25	≤ 0.125	≤ 0.125	0.06	0.25	0.125	0.5	0.5	0.125	0.125
0.55		0.12	0.12-8	4 – 16	0.12 – 0.25	8 – 32	2	8	0.5	32	32	2	2
0.97		0.0313	8	4	0.06 – 0.12	0.5 – 1	0.5	2	8	16	16	0.5	0.5
1.46		< 0.125 – 0.25	128	16	< 0.125 – 1	4	1 – 2	1 –> 16	1 –> 16	32 – 64	2 – 64	0.5 –> 16	2

142 Antifungal activity of Library A

143 All 13 compounds were tested against a panel of 4 strains from different clinically important
144 *Candida* species and an extended panel of *C. auris* (8 strains) and their antifungal activity and
145 lipophilicity was compared against the reference compound, Fluconazole (Table 2). The *C.*
146 *auris* panel covers a range of azole susceptibility profiles and clades (Clade I; TDG2211,
147 NCPF8971, Presumptive Clade I; TDG2512, Clade II; TDG2506, NCPF8984, Clade III; TDG1102,
148 NCPF8977, TDG1912). Whilst there is no defined EUCAST or CLSI resistance breakpoint for
149 fluconazole in *C. auris*, there is a presumed breakpoint of 32 µg/mL, therefore the panel
150 contains both azole 'susceptible' and azole 'resistant' isolates.

151

152 The goals of the structural modification was to understand the role of Fluorine and methyl at
153 the 4-position of the terminal aromatic or heteroaromatic ring, and the effect of introduction
154 of heteroatoms to the phenyl ring on antifungal activity of these compounds. Compound **1**
155 showed excellent activity across all *Candida* strains including the drug-resistant *C. auris* strains.
156 The MICs against all *Candida* strains was between 0.03 to 1 µg/mL, compared to fluconazole
157 which was inactive against the majority of the *C. auris* panel. Introduction of F at the 4-position
158 of the phenyl ring maintained activity across all strains with the exception of *C. auris* NCPF
159 8984 where a 4-fold reduction in activity was observed.

160

161 On the other hand, introducing a methyl group at the 4-position of the phenyl ring resulted in
162 significantly improved activity across all strains, including *C. auris* strains. In some cases, the
163 activity increased by as much as 20-fold. Introducing a heteroatom at the 2 position in the
164 form of nitrogen maintained activity across all strains, but a slight reduction was observed in
165 a few strains, particularly *N. glabrata* NCPF8018, where a 32-fold reduction was noted. A 4-
166 fold reduction was noted for *C. auris* TDG2506 and NCPF8984. Interestingly, the introduction
167 of a fluorine atom as seen in compound **5** restored activity to the levels observed for
168 compound **1** across all strains tested. This suggests that fluorine at the 4-position of the
169 pyridine ring is preferred over an unsubstituted pyridine, a preference not observed in the
170 case of the phenyl ring.

171

172 For the next set of compounds, the phenyl ring was replaced with a pyrimidine ring.
173 Compounds **6**, **7**, **8**, **9**, and **10** contained a pyrimidine ring with nitrogen atoms that are *ortho*-
174 to 2^o amine linkage. In the case of the unsubstituted pyrimidine (compound **6**), it maintained
175 excellent activity across all strains tested, with MICs ranging from 0.12 to 4 µg/mL. However,
176 for *C. auris* TDG2506 and NCPF8984, a 4-fold reduction was observed. Similar to what was
177 observed for compound **5**, where the introduction of fluorine restored activity, compound **7**,
178 which contains a 4-substituted fluorine, also showed a significant improvement in activity,
179 with MIC values comparable or in some cases better than compound **1**. Interestingly, only for
180 *C. auris* strain TDG1102 was an 8-fold reduction in activity observed. In contrast, for other
181 strains like TDG2506 and NCPF8984, a 30-fold improvement in activity was noted.

182

183 When the fluorine was replaced by a methyl group, as in compound **8**, activity was reduced,
184 particularly across all *C. auris* strains. This was surprising, given that in the case of compound
185 **3**, which contains a phenyl ring, replacing fluorine with methyl resulted in improved activity.
186 This suggests that the electronic environment of the terminal heteroaromatic ring, in this case,
187 pyrimidine, played a key role in the interaction of these compounds with the target enzyme,
188 affecting their activity.

189 In compound **9**, a dimethyl substitution was made at the 3 and 5 positions of the pyrimidine
190 ring. Surprisingly, this compound showed a significant improvement in activity, making it one
191 of the most active compounds, with activity comparable to that observed for compounds **1**
192 and **5**. Introducing a methyl substitution at the 3-position of compound **7**, as seen in
193 compound **10**, maintained activity, suggesting that these hydrophobic substitutions are well
194 tolerated in pyrimidine ring-containing compounds where nitrogen atoms are *ortho*- to 2^o
195 amine linkage.

196 Finally, for the Library A compounds, three further compounds were synthesized, where the
197 positions of the nitrogens in the pyrimidine ring were flipped compared to the previous series,
198 placing the nitrogens at *meta*- to 2^o amine linkage. The unsubstituted compound **11** showed
199 relatively poor activity compared to compound **1** (the phenyl compound), particularly against
200 *C. auris* strains, where reductions in activity of 32 to 64-fold were observed. Introducing
201 fluorine at the 4 position of this compound restored some activity, but it remained significantly
202 lower compared to both the unsubstituted phenyl ring compound **1** and the unsubstituted
203 pyrimidine compound **6**. Finally, when a methyl substitution was made at the 4 position of this
204 ring, as seen in compound **12**, a further reduction in MIC was observed. The structure-activity
205 relationship observed for the Library A compounds suggests that the effects of fluorine and
206 methyl substitutions on activity are highly dependent on the core aromatic or heteroaromatic
207 ring. These substitutions are well tolerated in phenyl and pyridyl rings, while their effects on
208 pyrimidine rings are more variable.

209

210

211 **Synthesis of the library B**

212 One interesting observation was the hydrophobicity, as measured by LogP, of the members
213 of Library A. Except for compounds **11** and **12**, all other compounds had a higher LogP,
214 indicating they are more hydrophobic compared to fluconazole (Table 2). Almost all of the
215 compounds, except for compound **13**, showed significantly superior activity compared to
216 fluconazole. This suggests that the hydrophobicity of theazole compounds potentially plays
217 an important role in their interaction with the target enzyme Erg11 and in the *in vitro*
218 activity of this compound series againstazole resistant *Candida* strains.

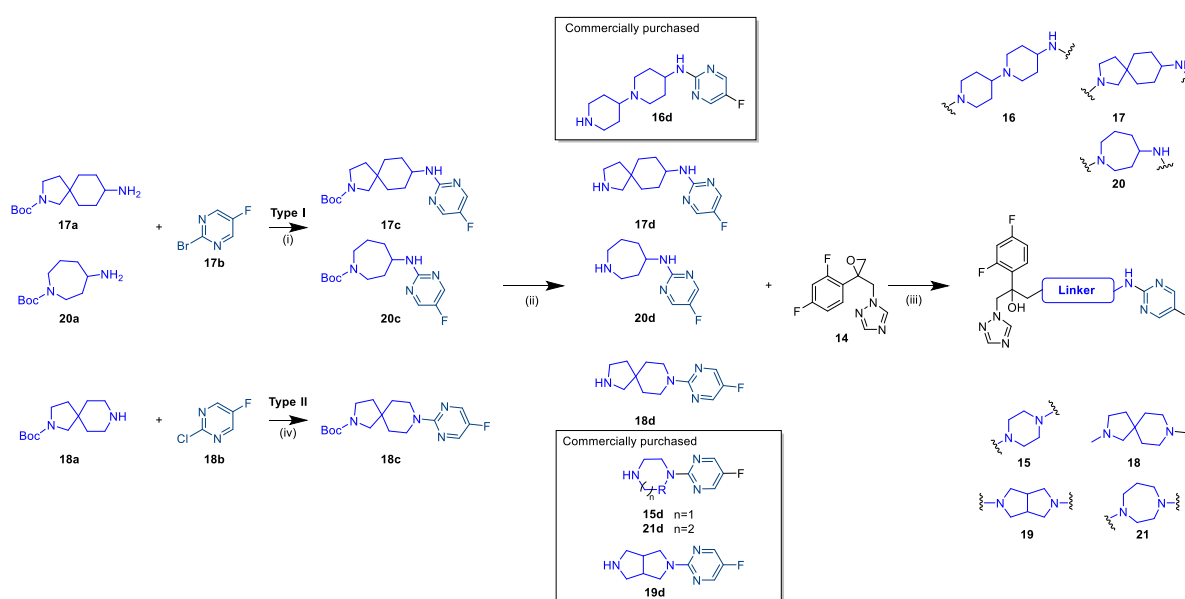
219 After reviewing the MIC activity data of Library A compounds, compound **7**, which contains a
220 pyrimidine ring with nitrogens at the *ortho* positions relative to the amine linker and a
221 fluorine atom at the 4-position, appeared to have the best overall activity profile. Therefore,

222 the terminal heteroaromatic fragment present in compound **7** was selected to design the
 223 next set of compounds, where the linker for aminopiperidine was varied. This was done
 224 specifically to explore the potential of introducing different types of linkers in the design and
 225 development of modified azole compounds that are active against resistant *Candida*
 226 strains—a strategy that has not been previously reported in the literature.

227 We selected seven different ring structures to modify compound **7**. These included various
 228 types of linkers, such as spiro and fused linkers, as well as six- and eight-membered linkers to
 229 introduce different types of ring strains. For compound **15**, we selected a piperazine linker. A
 230 [1,4'-bipiperidin]-4-amine was used as the linker for compound **16**, while 2-
 231 azaspiro[4.5]decan-8-amine was chosen for compound **17**. For compound **18**, we used 2,8-
 232 diazaspiro[4.5]decan-8-amine as the linker. A fused ring like octahydropyrrolo[3,4-c]pyrrole was
 233 selected for compound **19**, azepan-4-amine was chosen for compound **20**, and finally, a 1,4-
 234 diazepane ring, which is a seven-membered ring, was selected for compound **21**.

235 The linker-heteroaromatic fragments, **15a** to **21a** were either commercially obtained or
 236 synthesised according to Scheme 2. The final modified azole compounds **15** to **21** were
 237 synthesised following Scheme 2.

238



242 **Scheme 2.** General synthetic routes for library B, where terminal heteroaromatic fragment of
 243 **7** was connected to fluconazole core via different linkers. Conditions: (i) NaOtBu, JohnPhos,
 244 Pd₂(dba)₃, dioxane, 80 °C overnight; (ii) 4 M HCl in 4-dioxane, 2 h; (iii) Et₃N, EtOH, 80 °C,
 245 overnight; (iv) K₂CO₃, DMSO, 70 °C, overnight or EtN(Pr-i)₂, DMF, 130 °C, overnight;

246

247

248 Antifungal activity of library B

249 The synthesized compounds were evaluated against the same Candida panel, which included
250 clinically important Candida strains and drug-resistant *C. auris* strains. From the activity data
251 shown in Table 3, it can be seen that all linkers, except for the bipiperidine-4-amine linker
252 used in compound **16** and the fused linker octahydropyrrolo[3,4-c]pyrrole used in compound
253 **19**, were generally well tolerated. Compounds containing other linkers either maintained
254 their activity against these Candida strains, or in some cases, like compound **18**, showed
255 improved activity compared to compound **7** against most strains tested.

256

257 Compound **18**, which contains a 2,8-diazaspiro[4.5]decane linker demonstrated a 4- to 8-
258 fold improvement in activity across various Candida strains and had the best overall activity
259 profile among all the compounds synthesized. The compound was highly active against *C.*
260 *albicans* NCPF3281 (MIC ≤ 0.008 $\mu\text{g}/\text{mL}$), *C. parapsilosis* NCPF3209 (MIC 0.004 $\mu\text{g}/\text{mL}$) and
261 *C. auris* NCPF8977 (0.002 $\mu\text{g}/\text{mL}$). The MIC of this compound was less than 0.5 $\mu\text{g}/\text{mL}$
262 against all other strains. However, compound **15**, which contained a piperidine linker, also
263 showed good activity across the board, except against *C. auris* NCPF8984, where an
264 approximately 256-fold reduction in activity was observed and *C. auris* TDG2506 with a 64
265 fold reduction in activity. The compound had an MIC of 0.5 $\mu\text{g}/\text{mL}$ or less against all other
266 strains. Similarly, compound **17** maintained activity for most strains, but generally exhibited
267 a 2- to 8-fold reduction in activity (Table 3).

268

269 Compound **20**, on the other hand, showed some variability in activity. Its activity against *C.*
270 *tropicalis* NCPF8760 was 16-fold lower compared to compound **7**. Interestingly, against two
271 *C. auris* strains, TDG2211 and NCPF8984, compound **20** also showed significantly reduced
272 activity, with a more than 512-fold reduction observed. However, it showed comparable
273 activity against all other Candida strains.

274

275 Compound **21**, which had a 7-membered diazepane ring, generally maintained good activity
276 across all Candida strains tested. However, the activity was lower against *C. auris* strains,
277 with a 4- to 64-fold reduction observed for most strains, except for TDG2506, which showed
278 a more than 256-fold reduction in activity. It showed comparable activity against other
279 clinically important Candida strains with MICs ranging from 0.015 to 4 $\mu\text{g}/\text{mL}$ (Table 3).









280

281 From the activity profile of these second-generation compounds, it appears that linker
282 length did not play a specific role on activity. Compounds **16**, **17**, **18**, and **19**, which had
283 either fused or spiro linkers, generally had longer linkers compared to compounds **7**, **15**, **20**,
284 and **21**, but these lengths did not correlate with activity. For example, compound **18**, with a
285 longer spiro linker, and compound **7**, with a relatively short 4-aminopiperidine linker showed
286 the overall best activity profile and can be considered as lead compounds. These two

287 compounds showed a more than 128- to 512-fold increase in activity compared to
 288 fluconazole against *C. auris* strains. This suggests that the nature of the linker is more
 289 important than its length.

290

291 **Table 3.** Antifungal activity ($\mu\text{g/mL}$) of modified -fluconazole compounds against *Candida* from
 292 library B where different linker types were utilised.

Strains	Fluconazole	 7	 15	 16	 17	 18	 19	 20	 21
<i>C. albicans</i> NCPF3281	0.06– 0.125	≤ 0.12 5	≤ 0.01 6	0.015	0.031 3	≤ 0.008	0.12– 0.25	0.06– 0.125	≤ 0.016
<i>C. glabrata</i> NCPF8018	4	0.25	0.125	64	2	0.25– 0.5	1–2	1	0.25– 0.5
<i>C. tropicalis</i> NCPF8760	16	0.25	0.5	64	2	0.01– 0.5	8	4	1–2
<i>C. parapsilosis</i> NCPF3209	0.25	≤ 0.12 5	0.008	0.125	0.03	0.004– 0.008	0.125	0.12– 0.25	0.016– 0.06
<i>C. auris</i> TDG1912	>128	0.25	0.06– 0.12	4	0.5	0.12– 0.25	8–16	1	2
<i>C. auris</i> TDG2512	8	0.06	0.25	0.5	0.06– 0.12	0.03	4	0.25– 0.5	2
<i>C. auris</i> TDG1102	64	4	0.5	4	2	0.25	8–16	2	1–2
<i>C. auris</i> TDG2211	64	0.5	0.5	64	8	0.25	4	0.5	1
<i>C. auris</i> TDG2506	128	≤ 0.03	2	8	2	0.5	>128	16	8
<i>C. auris</i> NCPF8984	128	≤ 0.03	8	0.5	2	0.125	32	16	0.125
<i>C. auris</i> NCPF8971	32	0.25	0.125	0.5	0.25	0.125	1	0.25	0.125
<i>C. auris</i> NCPF8977	16	≤ 0.03	0.03	1	0.5	0.002– 0.03	1–4	0.25	0.12– 0.25

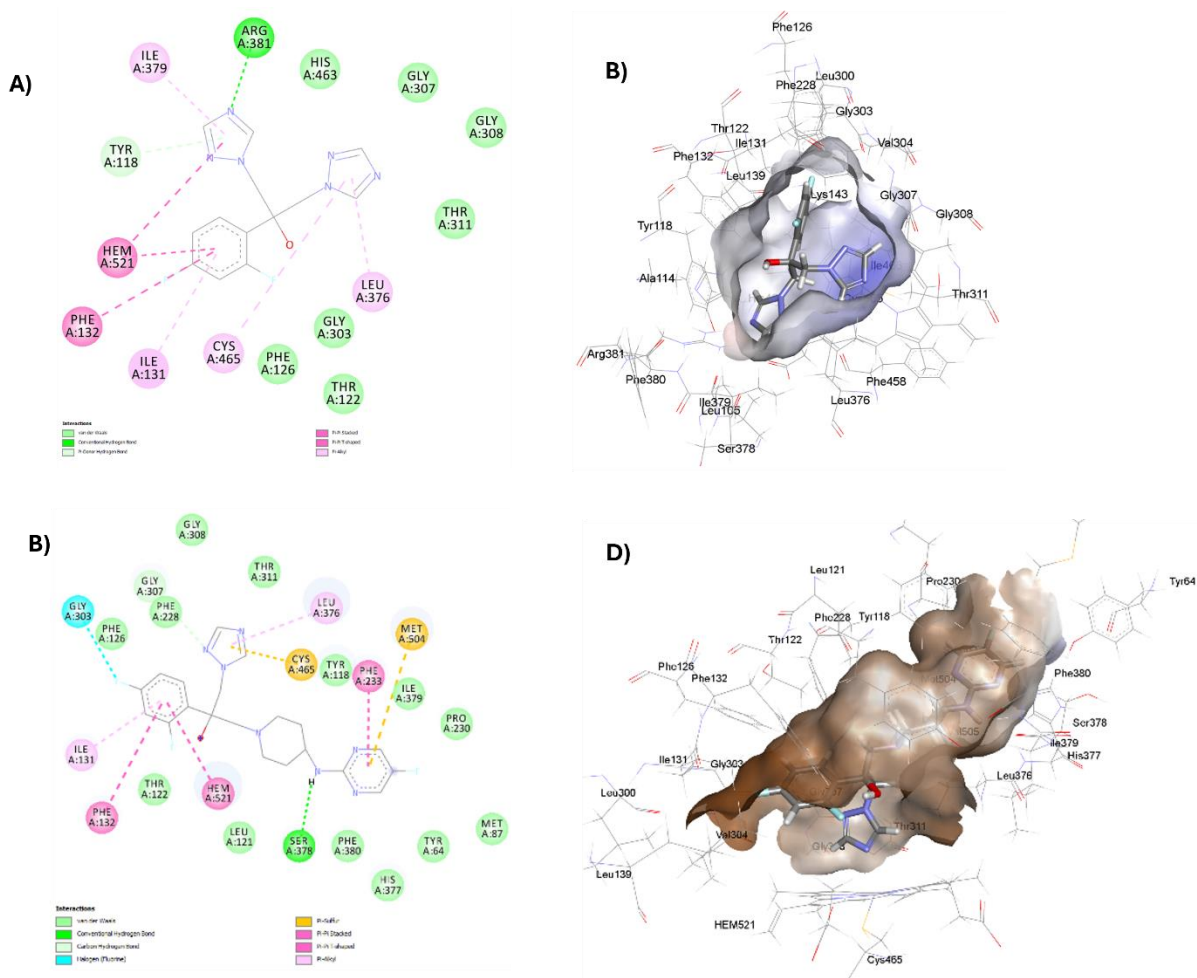
293

294

295 To further explore the differences in activity observed between fluconazole and the modified
 296 compound 7, particularly in *C. auris* strains, we evaluated the binding of these two compounds
 297 to the lanosterol 14 α -demethylase (LDM) enzyme in *C. auris* using molecular docking. As
 298 shown in Figure 2, interesting differences were observed in the 2D and 3D binding of both
 299 compounds. Fluconazole and compound 7 were found to bind to adjacent binding pockets.
 300 However, while fluconazole interacted with a more hydrophilic overall binding pocket,
 301 compound 7 interacted with a predominantly hydrophobic pocket and engaged with more

302 amino acid residues within the binding site. The length of the molecule may have contributed
 303 to these interactions, but the nature of the binding and the differences observed between
 304 compound 7 and fluconazole likely played a role in their ability to inhibit the target enzyme,
 305 and, their effectiveness in killing *C. auris* strains. The Chemscore and binding affinity values
 306 (Table 1) suggest that compound 7 binds more tightly to the LDM enzyme, and this tight
 307 binding, along with the differences in interactions, particularly in the hydrophobicity of key
 308 binding residues, may explain compound 7's superior ability to kill *C. auris* strains.

309



310

311

312 **Figure 2:** Differences in binding interactions observed for fluconazole (A & B), and compound
 313 7 (C & D) with lanosterol 14 α -demethylase (LDM) enzyme in *C. auris*

314

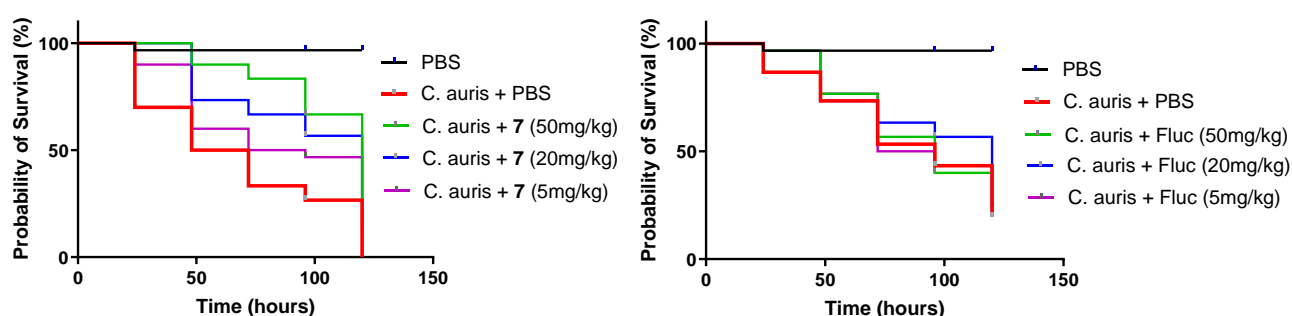
315

316 **Efficacy study in *G. mellonella* model**

317 After showing excellent *in vitro* activity against drug resistant *C. auris* strains, compound 7
 318 was selected to test the *in vivo* efficacy of these modified azoles in a *G. mellonella* model⁴⁴
 319 infected with *C. auris* (TDG1912). At first, the toxicity of the compound was tested using this

320 non-animal model, and compound **7** did not show any toxicity at 50 mg/Kg dose level
321 suggesting this modified azole scaffold is non-toxic at the doses studied. Compound **7**
322 expressed good protection over the infected larvae compared to fluconazole, which failed to
323 show any protection. The protection from the fungal infection recorded with the doses of 50
324 mg/kg and 20 mg/kg of **7** showed statistically significant protection (<0.0001 and 0.0025 with
325 Mantel-Cox log-rank test, respectively). Even at a low dose of 5 mg/kg, treated larvae showed
326 a reasonable survival rate of around 50% after 5 days of infection. As the dosage of **7**
327 increased, there was a progressive enhancement in larval survival rates. At the highest tested
328 dosage of 50 mg/kg, the final protection rate exceeded 70%, surpassing the approximately
329 75% larval mortality rate in the untreated group (Figure 3).

330



331

332 **Figure 3.** Efficacy of modified fluconazole compound **7** and fluconazole against *C. auris*
333 TDG1912 infection in *G. mellonella* model.

334

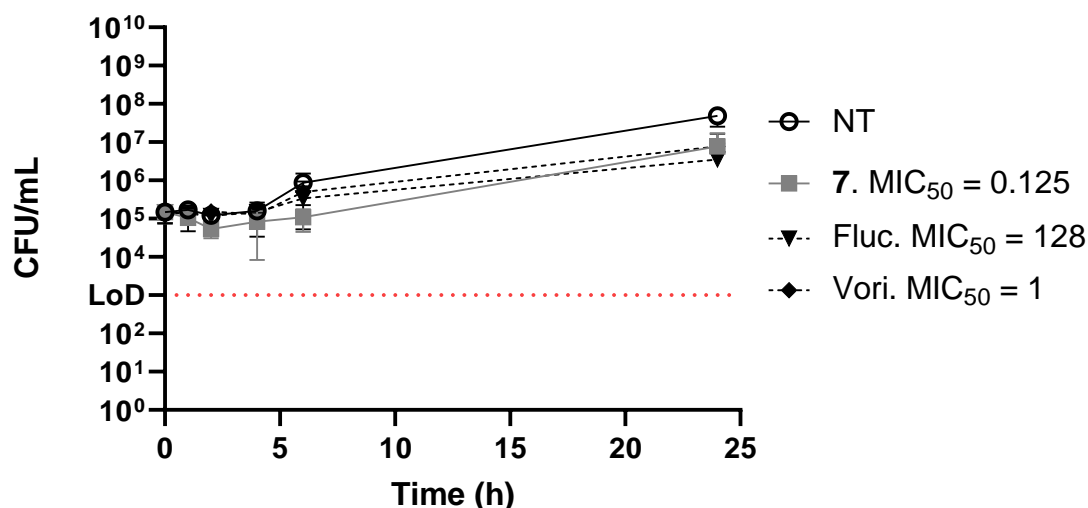
335 Time-kill assay of Compound **7**

336 Time-kill analysis, a dynamic assay using static MIC data, evaluates the kinetics and efficacy of
337 antifungal compounds over time. It monitors fungal growth at specific intervals after
338 treatment with antifungals at concentrations relative to the MIC. This method reveals
339 differences in the behavior of compounds with identical MICs, providing insights into their
340 mode of action. A reduction of colonies by ≤ 3 -log indicates fungistatic activity, while > 3 -log
341 reduction suggests fungicidal action. Azoles are typically fungistatic, but the higher *in vitro*
342 activity observed in modified azoles warranted an evaluation of their mode of action using the
343 time-kill kinetics assay. In this study, Compound **7**, fluconazole, and voriconazole were tested
344 against *C. auris* TDG1912 over 24 hours at 4x MIC₅₀ (Figure 4).

345 Compared to the untreated control sample, time-kill curves for all drug-treated samples
346 exhibited a slightly smaller fungal population size. Notably, compound **7** stood out among the
347 three tested compounds, demonstrating inhibition of fungal growth for at least six hours of
348 incubation. However, regrowth of *C. auris* TDG1912 was observed after the 6-hour time point.
349 It is important to note that the MIC value used for the time-kill assay corresponds to drug
350 concentrations inhibiting 50% of fungal growth, as defined by EUCAST guidelines for azoles.

351 All compounds, whether modified or commercially available, exhibited a fungistatic nature,
352 inhibiting fungal growth at varying efficacies, but not killing the fungi (Figure 4).

353



354

355 **Figure 4.** Time-kill of compound **7**, alongside two commercial reference antifungal drugs,
356 fluconazole (Fluc) and voriconazole (Vori) against *C. auris* TDG1912. The limit of the
357 detection (LoD) is 1000 CFU/mL. MIC₅₀ results are in µg/mL.

358

359 Conclusions

360 In summary, this study focussed on the synthesis and evaluation of a novel library of modified
361 azole compounds. The initial approach involved the use of a six-membered cyclic piperidine
362 linker, followed by the exploration of various heteroaliphatic linkers. Systematic investigation
363 of substitution patterns on the terminal aromatic or heteroaromatic fragment identified key
364 structural features that enhance antifungal activity against drug-resistant *C. auris* strains,
365 notably the impact of hydrophobicity, presence of heteroatom within the ring structure and
366 para-positioned substitutions on the aromatic ring. Compound **7** emerged as a promising
367 candidate, exhibiting potent activity against a broad panel of *Candida* strains. Further
368 exploration of different linkers helped establish the structure-activity relationship of this
369 modified azole series. The retention of activity across a wide range of linkers presents
370 opportunities for additional medicinal chemistry optimization during the lead optimization
371 phase. This research introduces a new azole-based chemical scaffold with enhanced activity
372 against all clinically important *Candida* strains, particularly drug-resistant *C. auris* strains.

373

374 Experimental Section

375 Chemistry

376 General Chemistry

377 ¹H NMR and ¹³C NMR spectra were recorded on a Bruker Spectrospin Spectrometer (400 MHz,
378 25 degrees) equipped with a SampleXpress autosampler system. Chemical shifts δH and δC
379 are recorded in parts per million (ppm) and referenced to the residual solvent peak. Coupling
380 constants (J) are recorded to the nearest 0.01 Hz. Assignment of ¹H and ¹³C NMR spectra was
381 made using the aid of TopSpin 3.5 software from Bruker, ACD/Labs or MestReNova from
382 Mestrelab Research. Purity is calculated using the HPLC-LCMS peak areas.

383 LC-MS system used is a Waters Alliance 2695 system with an elution in gradient. Low-
384 resolution mass spectra were analyzed and recorded on a Waters QZ instrument using
385 electrospray ionization (ESI) and coupled to a High-performance liquid chromatography
386 (HPLC) system. Selected mass-to-charge ratio peaks (m/z) are quoted in Daltons. HPLC grades
387 solvents were used for mobile phase and a Phenomenex Monolithic C18 50*4.60 mm column
388 was used for stationary phase. Detect method used UV detection performed on a Waters 2996
389 photo array detector.

390 Thin layer chromatography (TLC) was performed using Merck aluminium foil-backed sheets
391 precoated with 0.2 mm Kiieselgel 60 F254. Product spots were visualized by UV irradiation
392 (λ_{max} = 254 nm or 365 nm). Manual flash column chromatography was carried out using
393 Aldrich silica gel 60 Å, 40 – 63 μm (230 – 400 mesh). Eluting solvents and retention factors (R_f)
394 were indicated in the text.

395 The initial optical rotation of compounds was measured with a single wavelength polarimeter
396 ADP440 by Bellingham & Stanley at λ = 589 nm. Further optical rotatory dispersion (ORD)
397 measurements of selected compounds were measured with Chirascan spectrometers by
398 Applied Photophysics and were analysed with Pro-Data Chirascan.

399 All reactions were carried out in oven-dried glassware and all reagents were obtained
400 commercially either from Aldrich Chemicals Ltd., Alfa Aesar Ltd, Fluorochem, Fisher Scientific
401 or Apollo Chemicals Ltd.

402

403 **Chemical Synthesis**

404 ***General procedure for intermediate synthesis- (i) using reducing borohydride***

405 The aniline (1.0 equiv.) synthesised in the previous step or purchased commercially was
406 dissolved in the anhydrous DCM (0.1 mmol / mL) in the round-bottomed flask upon stirring,
407 followed by the addition of boc-protected linker fragment (1.5 equiv.), Na(OAc)₃BH (2.0 equiv.),
408 and AcOH (2.0 equiv.) successively. After the overnight stirring at ambient temperature, the
409 mixture was consecutively quenched with NaOH (aq.), extracted with DCM, dried with
410 anhydrous MgSO₄, and evaporated *in vacuo*. The crude mixture was purified with *flash* column
411 chromatography gradient-ly using chloroform and methanol to get the target pure product.

412 *tert-butyl 4-(phenylamino)piperidine-1-carboxylate 1a*

413

414 White solid (56%); ¹H NMR (400 MHz, 25 degree, CDCl₃) σ: 1.46 (9H, s), 2.04 (2H, dd, J = 13
415 Hz, 2 Hz), 2.44 (1H, t, J = 6 Hz), 2.88 (2H, t, J = 12 Hz), 3.42 (1H, tt, J = 10 Hz, 4 Hz), 3.72 (1H, t,

416 J = 6 Hz), 4.05 (2H, d, J = 8 Hz), 5.30 (1H, s), 6.68 – 6.81 (3H, m), 7.20 (2H, dd, J = 8 Hz, 7 Hz);
417 ¹³C NMR (101 MHz, 25 degree, CDCl₃) σ: 58.5, 32.0, 41.3, 79.8, 80.6, 115.3, 129.4, 129.6, 154.9,
418 208.0; m/z: (ESI⁺) 277.1 ([M+H]⁺); R_f: 0.9 (1 : 1 Hexane : Ethyl acetate); Purity: 96%.

419 *tert-butyl 4-(p-tolylamino)piperidine-1-carboxylate 3a*

420 Off-white solid (51%); ¹H NMR (400 MHz, 25 degree, CDCl₃) σ: 1.29 – 1.39 (2H, m), 1.46 (9H,
421 s), 2.03 (2H, dd, J = 12 Hz, 2 Hz), 2.24 (3H, s), 2.89 (2H, t, J = 12 Hz), 3.38 (1H, tt, J = 10 Hz, 4
422 Hz), 4.04 (2H, d, J = 9 Hz), 4.04 (1H, s, br), 6.60 (2H, d, J = 8 Hz), 7.00 (2H, d, J = 8 Hz); ¹³C NMR
423 (101 MHz, 25 degree, CDCl₃) σ: 20.5, 28.6, 32.3, 41.3, 51.4, 79.7, 114.6, 127.9, 130.0, 143.5,
424 154.9; m/z: (ESI⁺) 291.2 ([M+H]⁺); R_f: 0.03 (Ethyl acetate); Purity: ≥97%.

425 *tert-butyl 4-((2-fluoropyrimidin-5-yl)amino)piperidine-1-carboxylate 12a*

426 Off-white solid (49%); ¹H NMR (400 MHz, 25 degree, CDCl₃) σ: 1.41 (9H, s), 1.77 – 1.83 (1H,
427 m), 1.98 (2H, dd, J = 13 Hz, 3 Hz), 2.88 (2H, t, J = 12 Hz), 2.97 (1H, ddd, J = 13 Hz, 10 Hz, 3 Hz),
428 3.35 (1H, tt, J = 10 Hz, 4 Hz), 3.60 (1H, s, br), 4.03 (2H, d, J = 13 Hz), 7.97 (2H, s); ¹³C NMR (101
429 MHz, 25 degree, CDCl₃) σ: 28.4, 31.7, 34.2, 50.9, 67.6, 79.6, 80.0, 139.0 (d, J = 5 Hz), 144.7 (d,
430 J = 11 Hz), 154.7, 156.3 (d, J = 211 Hz); R_f: 0.2 (1 : 9 Ethyl acetate: DCM); Purity: ≥95%.

431 *tert-butyl 4-((2-methylpyrimidin-5-yl)amino)piperidine-1-carboxylate 13a*

432 White solid (53%); ¹H NMR (400 MHz, 25 degree, CDCl₃) σ: 1.36 – 1.43 (2H, m), 1.45 (9H, s),
433 2.01 (2H, dd, J = 13 Hz, 3 Hz), 2.65 (3H, s), 2.95 (2H, t, J = 12 Hz), 3.43 (1H, tt, J = 10 Hz, 4 Hz),
434 4.03 – 4.07 (2H, m), 4.42 (1H, s, br), 8.20 (2H, s); ¹³C NMR (101 MHz, 25 degree, CDCl₃) σ:
435 23.7, 28.6, 29.8, 31.9, 50.1, 80.0, 138.9, 141.1, 154.8, 154.8; m/z: (ESI⁺) 293.1 ([M+H]⁺); R_f:
436 0.5 (9 : 1 Ethyl acetate: MeOH); Purity: ≥98%.

437 **General procedure for intermediate synthesis- (ii) using base**

438 The starting material of pyrimidine (1.0 equiv.) in mixture of anhydrous MeCN (1.315 mmol /
439 mL) and dimethylacetamide (5.26 mmol / mL) was stirred under inert gas while BOC-
440 protected linker (1.2 equiv.) and DIPEA (1.2 equiv.) was added slowly. The reaction was carried
441 out in microwave at 160 °C for 1 h and at 180 °C for another 30 min. The crude mixture was
442 obtained by direct solvent evaporation. The mixture was purified by gradient flash column
443 chromatography in ethyl acetate and hexane with 1% TEA.

444 *5-methyl-N-(piperidin-4-yl)pyrimidin-2-amine 8a*

445 Pale-yellow solid (42%); ¹H NMR (400 MHz, 25 degree, MeOH-d₄) σ: 1.34 (2H, ddd, J = 24 Hz,
446 12 Hz, 3 Hz), 1.92 (2H, d, J = 12 Hz), 2.10 (3H, s), 2.92 (2H, t, J = 12 Hz), 3.02 (1H, tt, J = 11 Hz,
447 4 Hz), 4.65 (2H, d, J = 13 Hz), 8.16 (2H, s); ¹³C NMR (101 MHz, 25 degree, MeOH-d₄) σ: 14.4,
448 34.2, 44.0, 50.1, 119.7, 159.0, 151.6; m/z: (ESI⁺) 194.1 ([M+H]⁺); R_f : 0.1 (10 : 1 : 0.1 Ethyl
449 acetate : MeOH : TEA); Purity: ≥96%.

450 *tert-butyl 4-((4,6-dimethylpyrimidin-2-yl)amino)piperidine-1-carboxylate 9a*

451 Pale-yellow solid (45%); ¹H NMR (400 MHz, 25 degree, CDCl₃) σ : 1.20 – 1.30 (2H, m), 1.35
452 (9H, s), 1.91 (2H, d, J = 12 Hz), 2.15 (6H, s), 2.85 (2H, t, J = 12 Hz), 4.00 – 4.05 (1H, m), 4.59
453 (1H, d, J = 13 Hz), 6.18 (1H, s); ¹³C NMR (101 MHz, 25 degree, CDCl₃) σ : 23.7, 28.3, 32.2, 42.5,
454 47.5, 79.3, 109.6, 154.6, 161.4, 167.3; m/z: (ESI+) 307.2 ([M+H]⁺); R_f : 0.3 (2 : 8 Ethyl acetate
455 : Hexane); Purity: \geq 95%.

456 **General procedure for intermediate synthesis- (iii) using metal catalysis**

457 In an oven-dried round-bottomed flask containing anhydrous toluene (0.6 mmol / mL), the
458 pyrimidine fragment (1.0 equiv.), BOC-protected linker fragment (1.0 equiv.), rac-BINAP (0.04
459 equiv.), Pd₂(dba)₃ (0.02 equiv.) and KOtBu (1.2 equiv.) were added sequentially under inert
460 gas. The mixture was then heated under microwave at 100 °C for 2h. After cooling to room
461 temperature, the mixture was then filtered through celite. The filtrate was diluted with
462 saturated brine and extracted with ethyl acetate for three times before drying over anhydrous
463 MgSO₄ and evaporated in vacuo. Crude product was then purified with gradient flash column
464 chromatography in hexane and ethyl acetate.

465 *tert-butyl 4-((5-fluoro-4-methylpyrimidin-2-yl)amino)piperidine-1-carboxylate 10a*

466 Off-white solid (52%); ¹H NMR (400 MHz, 25 degree, CDCl₃) σ : 1.26 – 1.40 (2H, m), 1.44 (9H,
467 s), 1.99 (2H, d, J = 12 Hz), 2.31 (3H, d, J = 2 Hz), 2.92 (2H, t, J = 12 Hz), 3.83 – 4.01 (3H, m), 4.96
468 (1H, d, J = 7 Hz), 7.98 (1H, d, J = 1 Hz); ¹³C NMR (101 MHz, 25 degree, CDCl₃) σ : 17.8, 28.5,
469 32.2, 48.7, 79.7, 144.0 (d, J = 23 Hz), 151.1 (d, J = 246 Hz), 154.9, 155.8 (d, J = 16 Hz), 158.1 (d,
470 J = 3 Hz); m/z: (ESI+) 255.1 ([M – tBu + H]⁺); R_f 0.8 (Ethyl acetate); Purity: \geq 95%.

471 **Synthesis of fluconazole core**

472 *1-((2-(2,4-difluorophenyl)oxiran-2-yl)methyl)-1H-1,2,4-triazole (14)*

473 Trimethyl sulphoxonium iodide (2 equiv.) was added to toluene (0.12 mmol / mL) containing
474 1-(2,4-difluorophenyl)-2-(1H-1,2,4-triazol-1-yl)ethan-1-one (1 equiv.) and sodium hydroxide
475 30% (w/w) aqueous solution (10 equiv.). The mixture was heated under microwave (MW)
476 radiation for 50 minutes at 80 °C. Then the mixture was diluted with water and extracted with
477 ethyl acetate. The organic layer was combined and washed with saturated brine, dried over
478 anhydrous magnesium sulphate, and concentrated in vacuo. Pale yellow solid (52%); ¹H NMR
479 (400 MHz, 25 degree, CDCl₃) σ : 2.83 (1H, d, J = 5 Hz), 2.90 (1H, d, J = 5 Hz), 4.47 (1H, d, J = 15
480 Hz), 4.79 (1H, d, J = 15 Hz), 6.73 – 6.82 (2H, m), 7.11 – 7.16 (1H, m), 7.81 (1H, s), 8.03 (1H, s);
481 ¹³C NMR (101 MHz, 25 degree, CDCl₃) σ : 52.1, 53.4 (d, J = 3 Hz), 56.1 (d, J = 1 Hz), 104.0 (dd, J
482 = 26 Hz, J = 25 Hz), 111.6 (dd, J = 21 Hz, J = 3 Hz), 119.4 (dd, J = 15 Hz, 4 Hz), 129.5 (dd, J = 15
483 Hz, J = 10 Hz, 5 Hz), 144.0, 151.7, 160.5 (dd, J = 249 Hz, 12 Hz), 163.0 (dd, J = 251 Hz, 12 Hz);
484 m/z: (ESI+) 238.1 ([M+H]⁺); R_f: 0.24 (1 : 1 Ethyl acetate : Hexane); Purity: \geq 95%.

485

486 *tert-butyl 8-(5-fluoropyrimidin-2-yl)-2,8-diazaspiro[4.5]decane-2-carboxylate 18c*

487 Boc-protected linker fragment (1.0 equiv.) was dissolved in anhydrous DMF under inert gas
488 before the addition of pyrimidine fragment (1.275 equiv.) and DIPEA (3.125 equiv.). The

489 mixture was heated at 130 °C overnight and cooled to room temperature upon reaction
490 completion. After diluted with water, the mixture was extracted by ethyl acetate for three
491 times and dried over anhydrous MgSO₄ before evaporated *in vacuo*. The crude product was
492 purified with gradient *flash* column chromatography in mixture of DCM and ethyl acetate.
493 White solid (39%); ¹H NMR (400 MHz, 25 degree, CDCl₃) σ : 1.45 (9H, s), 1.58 (4H, dd, J = 11
494 Hz, 6 Hz), 1.75 (2H, t, J = 7 Hz), 3.19 (1H, s), 3.27 (1H, s), 3.38 – 3.43 (2H, m), 3.59 – 3.69 (2H,
495 m), 3.80 – 3.92 (2H, m), 8.16 (2H, s); ¹³C NMR (101 MHz, 25 degree, CDCl₃) σ : 28.7, 34.5, 36.5,
496 40.3, 42.1, 44.3 (d, J = 26 Hz), 55.6 (d, J = 75 Hz), 79.3, 145.2 (d, J = 22 Hz), 151.5 (d, J = 248
497 Hz), 154.9, 158.9; m/z: (ESI⁺) 281.1 ([M – tBu + H]⁺); R_f: 0.8 (1 : 1 DCM : Ethyl acetate); Purity:
498 \geq 95%.

499

500 *tert-butyl 4-((5-fluoropyrimidin-2-yl)amino)azepane-1-carboxylate 20c*

501

502 Synthesis was done by following the protocol described for **18c**. White solid (36%); ¹H NMR
503 (400 MHz, 25 degree, CDCl₃) σ : 1.46 (9H, s), 1.55 – 1.73 (3H, m), 1.83 – 1.99 (2H, m), 2.10 –
504 2.14 (1H, m), 3.24 (1H, tdd, J = 15 Hz, 9 Hz, 3 Hz), 3.39 – 3.48 (2H, m), 3.51 – 3.68 (1H, m), 3.94
505 (1H, s), 5.78 (1H, s, br), 8.20 (2H, s); ¹³C NMR (101 MHz, 25 degree, CDCl₃) σ : 24.5 (d, J = 16
506 Hz), 33.1 (d, J = 36 Hz), 35.0 (d, J = 4 Hz), 42.7 (d, J = 41 Hz), 52.08 (d, J = 24 Hz), 79.6, 105.5
507 (d, J = 8 Hz), 134.1 (d, J = 64 Hz), 145.5 (d, J = 23 Hz), 155.7; m/z: (ESI⁺) 311.2 ([M+H]⁺); R_f: 0.6
508 (98 : 2 Chloroform : MeOH); Purity: \geq 95%.

509

510 *tert-butyl 8-((5-fluoropyrimidin-2-yl)amino)-2-azaspiro[4.5]decane-2-carboxylate 17c*

511 Off-white solid (33%); ¹H NMR (400 MHz, 25 degree, CDCl₃) σ : 1.27 – 1.39 (2H, m), 1.45 (9H,
512 s), 1.61 – 1.68 (4H, m), 1.74 (1H, t, J = 7 Hz), 1.97 (2H, t, J = 13 Hz), 3.15 (2H, t, J = 32 Hz), 3.34
513 – 3.40 (2H, m), 3.77 (2H, m), 5.22 (1H, s, br), 8.18 (2H, s); m/z: (ESI⁺) 295.1 ([M – tBu + H]⁺); R_f
514 : 0.8 (1 : 1 DCM : Ethyl acetate); Purity: \geq 95%.

515

516 **General procedure for final compounds synthesis**

517 Fluconazole core (1 equiv.) and triethylamine (1.5 equiv.) was added into the solution of amine
518 (1.5 equiv.) in ethanol (0.07 mmol / mL) under stirring. The mixture was heated under stirring
519 at 80 oC overnight until all starting material disappeared. The solvent was removed under
520 reduced pressure and product was purified by flash column chromatography (MeOH in EtOAc,
521 0 – 20%).

522 *2-(2,4-difluorophenyl)-1-(4-(phenylamino)piperidin-1-yl)-3-(1H-1,2,4-triazol-1-
523 yl)propan-2-ol 1*

524 Light yellow liquid (58%); [α]_D24: 0 (c = 1.5 in methanol); ¹H NMR (400 MHz, 25 degree,
525 MeOH-d₄) σ : 1.37 – 1.47 (2H, m), 1.80 (1H, d, J = 15 Hz), 1.91 (1H, d, J = 11 Hz), 2.25 (1H, td, J
526 = 9 Hz, 3 Hz), 2.42 (1H, td, J = 2 Hz), 2.54 – 2.59 (1H, m), 2.73 – 2.78 (1H, m), 2.81 (1H, d, J =
527 13 Hz), 3.02 (1H, d, J = 14 Hz), 3.15 – 3.22 (1H, m), 4.58 (1H, d, J = 11 Hz), 4.68 (1H, d, J = 15
528 Hz), 6.56 – 6.61 (3H, M), 6.84 (1H, td, J = 9 Hz, 2 Hz), 6.90 – 6.96 (1H, m), 7.06 (2H, t, J = 8 Hz),
529 7.49 (1H, td, J = 8 Hz, 6 Hz), 7.75 (1H, s), 8.35 (1H, s); ¹³C NMR (101 MHz, 25 degree, MeOH-

530 d4) σ : 33.2, 50.8, 54.9, 57.6, 64.4, 74.5 (d, J = 6 Hz), 104.9 (dd, J = 28 Hz, 26 Hz), 114.9, 118.2,
531 127.5 (dd, J = 14 Hz, 4 Hz), 130.1, 130.8 (dd, J = 10 Hz, 6 Hz), 146.1, 149.0, 151.1, 152.0, 161.1
532 (dd, J = 246 Hz, 12 Hz), 164.2 (dd, J = 248 Hz, 12 Hz); m/z: (ESI+) 414.2 ([M+H]⁺); HRMS: (ESI+)
533 found 414.2088, ([M+H]⁺) requires 414.2100; Rf : 0.4 (Ethyl acetate); Purity: \geq 99%.

534 *2-(2,4-difluorophenyl)-1-(4-((4-fluorophenyl)amino)piperidin-1-yl)-3-(1H-1,2,4-triazol-*
535 *1-yl)propan-2-ol 2*

536 Light yellow solid (50%); M.P. = 93 – 102 °C; $[\alpha]_{D24}$: 14 (c = 0.7 in methanol); ¹H NMR (400
537 MHz, 25 degree, MeOH-d4) σ : 2.38 (3H, m), 2.47 – 2.48 (4H, m), 2.79 (1H, d, J = 14 Hz), 2.99
538 (1H, dd, J = 14 Hz, 1 Hz), 3.43 – 3.50 (2H, m), 4.59 (1H, d, J = 14 Hz), 4.68 (1H, d, J = 14 Hz),
539 6.84 (1H, td, J = 8 Hz, 3 Hz), 6.88 – 6.94 (1H, m), 6.97 (1H, td, J = 8 Hz, 2 Hz), 7.06 (1H, dt, J =
540 10 Hz, 2 Hz), 7.09 (1H, d, J = 8 Hz), 7.30 (1H, td, J = 8 Hz, 7 Hz), 7.46 (1H, td, J = 9 Hz, 7 Hz), 7.74
541 (1H, s), 8.33 (1H, s); ¹³C NMR (101 MHz, 25 degree, MeOH-d4) σ : 54.1, 55.4, 57.5 (d, J = 5 Hz),
542 63.2 (d, J = 2 Hz), 64.5 (d, J = 3 Hz), 74.9 (d, J = 4 Hz), 104.9 (td, J = 28 Hz, 2 Hz), 112.0 (dd, J =
543 21 Hz, 3 Hz), 115.0 (d, J = 19 Hz), 117.0 (d, J = 20 Hz), 126.3 (d, J = 3 Hz), 127.2 (dd, J = 12 Hz,
544 4 Hz), 130.8 (dd, J = 3 Hz, 1 Hz), 131.0 (d, J = 8 Hz), 141.5 (d, J = 7 Hz), 146.1, 151.1, 159.7 (d, J
545 = 28 Hz), 162.9 (dd, J = 246 Hz, 12 Hz), 164.3 (dd, J = 246 Hz, 12 Hz); m/z: (ESI+) 432.2 ([M+H]⁺);
546 HRMS: (ESI+) found 432.1998, ([M+H]⁺) requires 432.2006; Rf : 0.4 (Ethyl acetate); Purity:
547 \geq 98%.

548 *2-(2,4-difluorophenyl)-1-(4-(p-tolylamino)piperidin-1-yl)-3-(1H-1,2,4-triazol-1-*
549 *yl)propan-2-ol 3*

550 Transparent liquid (60%); $[\alpha]_{D26}$: -4 (c = 2.5 in methanol); ¹H NMR (400 MHz, 25 degree,
551 MeOH-d4) σ : 1.25 – 1.44 (2H, m), 1.76 – 1.81 (1H, m), 1.87 – 1.91 (1H, m), 2.17 (3H, s), 2.23
552 (1H, td, J = 12 Hz, 3 Hz), 2.40 (1H, td, J = 12 Hz, 3 Hz), 2.53 – 2.57 (1H, m), 2.72 – 2.76 (1H, m),
553 2.80 (1H, d, J = 14 Hz), 3.00 (1H, dd, J = 14 Hz, 2 Hz), 3.13 (1H, tt, J = 10 Hz, 4 Hz), 4.57 (1H, d,
554 J = 14 Hz), 4.67 (1H, d, J = 14 Hz), 6.55 (2H, ddd, J = 8 Hz, 3 Hz, 2 Hz), 6.84 (1H, tdd, J = 9 Hz, 3
555 Hz, 1 Hz), 6.89 – 6.95 (3H, m), 7.48 (1H, td, J = 9 Hz, 7 Hz), 7.74 (1H, s), 8.34 (1H, s); ¹³C NMR
556 (101 MHz, 25 degree, MeOH-d4) σ : 20.5, 33.2, 33.5, 51.4, 54.9, 55.7, 57.6 (d, J = 5 Hz), 64.3
557 (d, J = 4 Hz), 74.4 (d, J = 6 Hz), 104.9 (dd, J = 28 Hz, 26 Hz), 112.0 (dd, J = 21 Hz, 3 Hz), 115.6,
558 127.5 (dd, J = 13 Hz, 4 Hz), 127.8, 130.5, 130.8 (dd, J = 10 Hz, 6 Hz), 146.1, 146.4, 151.1, 160.7
559 (dd, J = 246 Hz, 12 Hz), 164.1 (dd, J = 248 Hz, 12 Hz); m/z: (ESI+) 428.2 ([M+H]⁺); HRMS: (ESI+)
560 found 428.2254, ([M+H]⁺) requires 428.22564; Rf : 0.4 (Ethyl acetate); Purity: \geq 98%.

561 *2-(2,4-difluorophenyl)-1-(4-(pyridin-2-ylamino)piperidin-1-yl)-3-(1H-1,2,4-triazol-1-*
562 *yl)propan-2-ol 4*

563 Light yellow solid (52%); M.P. = 54 – 65 °C; $[\alpha]_{D26}$: 0 (c = 1.2 in methanol); ¹H NMR (400 MHz,
564 25 degree, MeOH-d4) σ : 1.33 – 1.52 (2H, m), 1.79 (1H, d, J = 12 Hz), 1.90 (1H, d, J = 12 Hz),
565 2.28 (1H, td, J = 12 Hz, 1 Hz), 2.44 (1H, td, J = 13 Hz, 2 Hz), 2.57 (1H, d, J = 11 Hz), 2.75 (1H, d, J
566 = 10 Hz), 2.81 (1H, d, J = 15 Hz), 3.02 (1H, d, J = 14 Hz), 3.57 (1H, tt, J = 10 Hz, 3 Hz), 4.59 (1H,
567 d, J = 11 Hz), 4.68 (1H, d, J = 9 Hz), 6.46 – 6.50 (2H, m), 6.84 (1H, td, J = 7 Hz, 4 Hz), 6.92 (1H,
568 td, J = 10 Hz, 2 Hz), 7.37 (1H, td, J = 8 Hz, 3 Hz), 7.49 (1H, dd, J = 12 Hz, 8 Hz), 7.75 (1H, s), 7.86
569 (1H, d, J = 7 Hz), 8.35 (1H, s); ¹³C NMR (101 MHz, 25 degree, MeOH-d4) σ : 33.2, 33.4, 49.9,
570 54.9, 55.6, 57.6 (d, J = 7 Hz), 64.4 (d, J = 7 Hz), 74.5 (d, J = 8 Hz), 104.9 (dd, J = 28 Hz, J = 26 Hz),

571 110.4, 112.0 (dd, J = 21 Hz, 5 Hz), 113.0, 126.7 (dd, J = 17 Hz, 6 Hz), 130.8 (dd, J = 9 Hz, 5 Hz),
572 138.7, 146.1, 147.8, 151.1, 159.5, 160.7 (dd, J = 256 Hz, 12 Hz), 164.1 (dd, J = 248 Hz, 12 Hz);
573 m/z: (ESI+) 415.2 ([M+H]⁺); HRMS: (ESI+) found 415.2039, ([M+H]⁺) requires 415.2052; Rf :
574 0.4 (10 : 1 Ethyl acetate : MeOH); Purity: ≥95%.

575 *2-(2,4-difluorophenyl)-1-(4-((5-fluoropyridin-2-yl)amino)piperidin-1-yl)-3-(1H-1,2,4-*
576 *triazol-1-yl)propan-2-ol 5*

577 Light yellow solid (52%); M.P. = 59 – 74 °C; [α]_D24: 7.4 (c = 1.4 in methanol); (400 MHz, 25
578 degree, MeOH-d₄) σ: 1.31 – 1.50 (2H, m), 1.79 (1H, d, J = 17 Hz), 1.90 (1H, d, J = 17 Hz), 2.26
579 (1H, td, J = 8 Hz, 8 Hz), 2.42 (1H, td, J = 13 Hz, 3 Hz), 2.56 (1H, d, J = 19 Hz), 2.74 (1H, d, J = 14
580 Hz), 2.81 (1H, d, J = 20 Hz), 3.01 (1H, d, J = 16 Hz), 3.54 (1H, tt, J = 11 Hz, 4 Hz), 4.60 (1H, d, J =
581 15 Hz), 4.68 (1H, d, J = 14 Hz), 6.46 (1H, dd, J = 11 Hz, 5 Hz), 6.84 (1H, td, J = 10 Hz, 4 Hz), 6.92
582 (1H, td, J = 11 Hz, 3 Hz), 7.22 (1H, td, J = 11 Hz, 2 Hz), 7.48 (1H, dd, J = 15 Hz, 12 Hz), 7.75 (1H,
583 s), 7.76 (1H, m), 8.35 (1H, s); ¹³C NMR (101 MHz, 25 degree, MeOH-d₄) σ: 33.1, 33.4, 49.9,
584 54.9, 55.6, 57.6 (d, J = 7 Hz), 64.4 (d, J = 6 Hz), 74.5 (d, J = 5 Hz), 104.9 (dd, J = 26 Hz, J = 23
585 Hz), 110.8 (d, J = 4 Hz), 112.0 (dd, J = 21 Hz, 5 Hz), 126.7 (d, J = 22 Hz), 127.4 (dd, J = 11 Hz, 6
586 Hz), 130.8 (dd, J = 12 Hz, 7 Hz), 134.0 (d, J = 19 Hz), 146.1, 151.1, 154.2 (d, J = 249 Hz), 156.7,
587 160.7 (dd, J = 253 Hz, 12 Hz), 164.1 (dd, J = 246 Hz, 12 Hz); m/z: (ESI+) 433.1 ([M+H]⁺); HRMS:
588 (ESI+) found 433.1948, ([M+H]⁺) requires 433.1958; Rf : 0.1 (Ethyl acetate); Purity: ≥95%.

589 *2-(2,4-difluorophenyl)-1-(4-(pyrimidin-2-ylamino)piperidin-1-yl)-3-(1H-1,2,4-triazol-1-*
590 *yl)propan-2-ol 6*

591 Light orange solid (52%); M.P. = 134 – 139 °C; [α]_D24: 20 (c = 1.0 in methanol); ¹H NMR (400
592 MHz, 25 degree, MeOH-d₄) σ: 1.39 – 1.58 (2H, m), 1.84 (2H, dd, J = 49 Hz, 11 Hz), 2.26 (1H,
593 td, J = 10 Hz, 5 Hz), 2.44 (1H, td, J = 11 Hz, 4 Hz), 2.58 (1H, d, J = 8 Hz), 2.77 (1H, d, J = 9 Hz),
594 2.81 (1H, d, J = 19 Hz), 3.02 (1H, d, J = 22 Hz), 3.69 (1H, tt, J = 12 Hz, 3 Hz), 4.60 (1H, d, J = 9
595 Hz), 4.69 (1H, d, J = 16 Hz), 6.55 (1H, t, J = 4 Hz), 6.85 (1H, td, J = 12 Hz, 3 Hz), 6.93 (1H, td, J =
596 10 Hz, 3 Hz), 7.49 (1H, dd, J = 16 Hz, 8 Hz), 7.75 (1H, s), 8.21 (1H, s), 8.23 (1H, s), 8.36 (1H, s);
597 ¹³C NMR (101 MHz, 25 degree, MeOH-d₄) σ: 33.1, 32.9, 49.9, 54.9, 55.6, 57.6 (d, J = 5 Hz),
598 64.4 (d, J = 8 Hz), 74.6 (d, J = 7 Hz), 104.9 (dd, J = 33 Hz, 17 Hz), 111.2, 112.0 (dd, J = 24 Hz, 6
599 Hz), 127.4 (dd, J = 12 Hz, J = 4 Hz), 130.8 (dd, J = 10 Hz, 3 Hz), 146.1, 151.1, 159.3, 160.7 (dd,
600 J = 250 Hz, 12 Hz), 162.85, 164.14 (dd, J = 248 Hz, 12 Hz); m/z: (ESI+) 416.2 ([M+H]⁺); HRMS:
601 (ESI+) found 416.1994, ([M+H]⁺) requires 416.2005; Rf : 0.4 (9 : 1 Ethyl acetate : MeOH);
602 Purity: ≥95%.

603 *2-(2,4-difluorophenyl)-1-(4-((5-fluoropyrimidin-2-yl)amino)piperidin-1-yl)-3-(1H-1,2,4-*
604 *triazol-1-yl)propan-2-ol 7*

605 Pale yellow crystals (58%); M.P. = 119 °C; [α]_D23: 24 (c = 0.9 in methanol); ¹H NMR (400 MHz,
606 25 degree, MeOH-d₄) σ: 1.38 – 1.56 (2H, m), 1.79 (1H, d, J = 12 Hz), 1.90 (1H, d, J = 11 Hz),
607 2.25 (1H, td, J = 12 Hz, 1 Hz), 2.42 (1H, td, J = 12 Hz, 2 Hz), 2.57 (1H, d, J = 9 Hz), 2.75 (1H, d, J
608 = 9 Hz), 2.81 (1H, d, J = 16 Hz), 3.01 (1H, d, J = 10 Hz), 3.64 (1H, tt, J = 9 Hz, 4 Hz), 4.59 (1H, d,
609 J = 11 Hz), 4.68 (1H, d, J = 12 Hz), 6.82 – 6.96 (2H, m), 7.49 (1H, dd, J = 15 Hz, 8 Hz), 7.75 (1H,
610 s), 8.17 (1H, s), 8.35 (1H, s); ¹³C NMR (101 MHz, 25 degree, MeOH-d₄) σ: 32.8, 33.1, 49.3,
611 54.9, 55.6, 57.6 (d, J = 5 Hz), 64.3 (d, J = 3 Hz), 74.54 (d, J = 4 Hz), 104.9 (t, J = 24 Hz), 112.0

612 (dd, J = 21 Hz, 3 Hz), 127.4 (dd, J = 13 Hz, 4 Hz), 130.8 (dd, J = 8 Hz, 5 Hz), 146.6 (d, J = 21 Hz),
613 151.1, 152.0, 154.4, 160.4 (d, J = 1 Hz), 164.2 (dd, J = 246 Hz, 12 Hz), 160.7 (dd, J = 243 Hz, 12
614 Hz); m/z: (ESI+) 434.2 ([M+H]⁺); HRMS: (ESI+) found 434.1902, ([M+H]⁺) requires 434.1911;
615 Rf : 0.2 (Ethyl acetate); Purity: ≥98%.

616 *2-(2,4-Difluorophenyl)-1-(4-((5-methylpyrimidin-2-yl)amino)piperidin-1-yl)-3-(1H-*
617 *1,2,4-triazol-1-yl)propan-2-ol 8*

618 Transparent liquid (51%); [α]_D24: 0 (c = 1.5 in methanol); ¹H NMR (400 MHz, 25 degree,
619 methanol-d₄) σ: 1.23 (2H, t, J = 10 Hz), 1.87 (2H, t, J = 13 Hz), 2.12 (3H, s), 2.62 (1H, tt, J = 9
620 Hz, 4 Hz), 2.92 (1H, t, J = 12 Hz), 3.08 (1H, d, J = 12 Hz), 3.22 (1H, d, J = 12 Hz), 4.52 (2H, s),
621 4.65 (1H, d, J = 14 Hz), 4.72 (1H, d, J = 14 Hz), 6.86 (1H, t, J = 9 Hz), 6.94 (1H, t, J = 10 Hz), 7.48
622 (1H, dd, J = 14 Hz, 8 Hz), 7.78 (1H, s), 8.16 (2H, s), 8.34 (1H, s); ¹³C NMR (101 MHz, 25 degree,
623 MeOH-d₄) σ: 17.6, 32.9, 33.2, 54.9, 55.6, 57.6 (d, J = 5 Hz), 64.3 (d, J = 4 Hz), 74.5 (d, J = 6 Hz),
624 104.9 (dd, J = 28 Hz, 26 Hz), 112.0 (dd, J = 21 Hz, 3 Hz), 127.4 (dd, J = 13 Hz, 4 Hz), 130.8 (dd, J
625 = 9 Hz, 6 Hz), 144.9 (d, J = 24 Hz), 146.1, 151.1, 151.9 (d, J = 244 Hz), 157.0 (d, J = 16 Hz), 160.7
626 (dd, J = 247 Hz, 12 Hz), 159.7 (d, J = 3 Hz), 164.1 (dd, J = 248 Hz, 12 Hz); m/z: (ESI+) 430.2
627 ([M+H]⁺); HRMS: (ESI+) found 430.2151, ([M+H]⁺) requires 430.2161; Rf : 0.4 (9 : 1 Ethyl
628 acetate : MeOH); Purity: ≥98%.

629 *2-(2,4-Difluorophenyl)-1-(4-((4,6-dimethylpyrimidin-2-yl)amino)piperidin-1-yl)-3-(1H-*
630 *1,2,4-triazol-1-yl)propan-2-ol 9*

631 Light orange solid (52%); M.P. = 160 – 162 °C; [α]_D24: -11 (c = 0.9 in methanol); ¹H NMR (400
632 MHz, 25 degree, methanol-d₄) σ: 1.36 – 1.56 (2H, m), 1.76 – 1.82 (1H, m), 1.86 – 1.93 (1H, m),
633 2.23 (6H, s), 2.28 (1H, td, J = 11.4 Hz, 3 Hz), 2.44 (1H, td, J = 12 Hz, 3 Hz), 2.53 – 2.59 (1H, m),
634 2.72 – 2.77 (1H, m), 2.81 (1H, d, J = 14 Hz), 3.02 (1H, dd, J = 14 Hz, 2 Hz), 3.78 (1H, tt, J = 11
635 Hz, 4 Hz), 4.59 (1H, d, J = 14 Hz), 4.69 (1H, d, J = 14 Hz), 6.37 (1H, s), 6.85 (1H, tdd, J = 8 Hz, 3
636 Hz, 1 Hz), 6.93 (1H, ddd, J = 12 Hz, 9 Hz, 3 Hz), 7.49 (1H, td, J = 9 Hz, 7 Hz), 7.75 (1H, s), 8.36
637 (1H, s); ¹³C NMR (101 MHz, 25 degree, MeOH-d₄) σ: 23.6, 33.1, 33.4, 54.9, 55.6, 57.6 (d, J =
638 5 Hz), 64.4 (d, J = 4 Hz), 75.6 (d, J = 6 Hz), 104.9 (dd, J = 28 Hz, 26 Hz), 110.3, 112.0 (dd, J = 21
639 Hz, 3 Hz), 127.4 (dd, J = 13 Hz, 4 Hz), 130.8 (dd, J = 9 Hz, 6 Hz), 146.1, 151.1, 160.7 (dd, J = 246
640 Hz, 12 Hz), 162.9, 164.2 (dd, J = 247 Hz, 12 Hz), 169.1; m/z: (ESI+) 444.2 ([M+H]⁺); HRMS:
641 (ESI+) found 444.2311, ([M+H]⁺) requires 444.2318; Rf : 0.5 (9 : 1 Ethyl acetate : MeOH);
642 Purity: ≥98%.

643 *2-(2,4-Difluorophenyl)-1-(4-((5-fluoro-4-methylpyrimidin-2-yl)amino)piperidin-1-yl)-3-*
644 *(1H-1,2,4-triazol-1-yl)propan-2-ol 10*

645 White solid (52%); M.P. = 79 – 130 °C; [α]_D24: 17 (c = 0.6 in methanol); ¹H NMR (400 MHz, 25
646 degree, methanol-d₄) σ: 1.38 – 1.56 (2H, m), 1.80 (1H, d, J = 12 Hz), 1.90 (1H, d, J = 12 Hz),
647 2.26 (1H, t, J = 11 Hz), 2.30 (3H, s), 2.43 (1H, t, J = 11 Hz), 2.58 (1H, d, J = 11 Hz), 2.76 (1H, d, J
648 = 11 Hz), 2.82 (1H, d, J = 13 Hz), 3.03 (1H, d, J = 13 Hz), 3.66 (1H, tt, J = 10 Hz, 4 Hz), 4.60 (1H,
649 d, J = 14 Hz), 4.70 (1H, d, J = 14 Hz), 6.86 (1H, td, J = 7 Hz, 2 Hz), 6.94 (1H, td, J = 9 Hz, 2 Hz),
650 7.50 (1H, dd, J = 16 Hz, 9 Hz), 7.77 (1H, s), 8.01 (1H, s), 8.37 (1H, s); ¹³C NMR (101 MHz, 25
651 degree, MeOH-d₄) σ: 17.6, 32.9, 33.2, 54.9, 55.6, 57.6 (d, J = 5 Hz), 64.3 (d, J = 4 Hz), 74.5 (d,
652 J = 6 Hz), 104.9 (dd, J = 28 Hz, 26 Hz), 112.0 (dd, J = 21 Hz, 3 Hz), 127.4 (dd, J = 13 Hz, 3 Hz),

653 130.8 (dd, J = 9 Hz, 6 Hz), 145.0 (d, J = 24 Hz), 146.1, 151.1, 151.9 (d, J = 244 Hz), 157.0 (d, J =
654 15 Hz), 159.7 (d, J = 3 Hz), 160.7 (dd, J = 246 Hz, 12 Hz), 164.2 (dd, J = 248 Hz, 12 Hz); m/z:
655 (ESI+) 448.2 ([M+H]⁺); HRMS: (ESI+) found 448.2057, ([M+H]⁺) requires 448.2067; Rf : 0.3
656 (Ethyl acetate); Purity: ≥97%.

657 *2-(2,4-difluorophenyl)-1-(4-(pyrimidin-5-ylamino)piperidin-1-yl)-3-(1H-1,2,4-triazol-1-*
658 *yl)propan-2-ol 11*

659 Light yellowish solid (49%); M.P. = 141 – 198 °C; [α]_D26: 17 (c = 0.6 in methanol); ¹H NMR
660 (400 MHz, 25 degree, MeOH-d₄) σ: 1.38 – 1.50 (2H, m), 1.87 (2H, dt, J = 37 Hz, 2 Hz), 2.31 (1H,
661 td, J = 12 Hz, 4 Hz), 2.45 (1H, td, J = 12 Hz, 3 Hz), 2.62 (1H, d, J = 11 Hz), 2.78 (1H, H, J = 12 Hz),
662 2.83 (1H, d, J = 14 Hz), 3.04 (1H, dd, J = 12 Hz, 3 Hz), 3.29 (1H, tt, J = 12 Hz, 3 Hz), 4.60 (1H, d,
663 J = 10 Hz), 4.69 (1H, d, J = 14 Hz), 6.85 (1H, td, J = 9 Hz, 3 Hz), 6.93 (1H, td, J = 10 Hz, 3 Hz),
664 7.49 (1H, td, J = 9 Hz, 7 Hz), 7.76 (1H, s), 8.09 (1H, s), 8.32 (1H, s), 8.35 (1H, s); ¹³C NMR (101
665 MHz, 25 degree, MeOH-d₄) σ: 32.8, 54.5, 57.6, 64.3, 64.4, 74.7 (d, J = 5 Hz), 104.9 (dd, J = 28
666 Hz, 21 Hz), 112.0 (dd, J = 21 Hz, 4 Hz), 127.3 (dd, J = 13 Hz, 4 Hz), 130.8 (dd, J = 9 Hz, 6 Hz),
667 141.4, 143.7, 146.1, 146.8, 151.1, 160.7 (dd, J = 246 Hz, 12 Hz), 164.15 (dd, J = 245 Hz, 12 Hz);
668 m/z: (ESI+) 416.2 ([M+H]⁺); HRMS: (ESI+) found 416.1994, ([M+H]⁺) requires 416.2005; Rf :
669 0.1 (9 : 1 Ethyl acetate : MeOH); Purity: ≥95%.

670 *2-(2,4-difluorophenyl)-1-(4-((2-fluoropyrimidin-5-yl)amino)piperidin-1-yl)-3-(1H-1,2,4-*
671 *triazol-1-yl)propan-2-ol 12*

672 Light white liquid (54%); [α]_D25: 28 (c = 0.4 in methanol); ¹H NMR (400 MHz, 25 degree,
673 MeOH-d₄) σ: 1.66 – 1.92 (2H, m), 2.17 – 2.34 (1H, m), 2.42 (1H, td, J = 11 Hz, 3 Hz), 2.53 – 2.60
674 (1H, m), 2.65 – 2.82 (2H, m), 3.02 (1H, dd, J = 14 Hz, 1 Hz), 3.15 (1H, tt, J = 10 Hz, 4 Hz), 3.87
675 (2H, s), 4.60 (1H, d, J = 7 Hz), 4.66 (1H, d, J = 4 Hz), 6.85 (1H, td, J = 8 Hz, 2 Hz), 6.92 (1H, ddt,
676 J = 12 Hz, 9 Hz, 2 Hz), 7.49 (1H, dddd, J = 12 Hz, 7 Hz, 7 Hz, 3 Hz), 7.75 (1H, d, J = 4 Hz), 7.95
677 (2H, s), 8.35 (1H, s); ¹³C NMR (101 MHz, 25 degree, MeOH-d₄) σ: 32.9, 35.1, 35.3, 55.2, 57.6
678 (d, J = 5 Hz), 64.3 (d, J = 4 Hz), 74.6 (d, J = 6 Hz), 104.9 (dd, J = 28 Hz, 26 Hz), 112.0 (dd, J = 21
679 Hz, 3 Hz), 127.4 (dd, J = 9 Hz, 4 Hz), 130.8 (dd, J = 10 Hz, 6 Hz), 138.9, 145.1, 146.1, 151.1,
680 159.5, 160.7 (dd, J = 246 Hz, 12 Hz), 164.13 (dd, J = 248 Hz, 12 Hz); m/z: (ESI+) 446.2; Rf : 0.3
681 (9 : 1 Ethyl acetate : MeOH); Purity: ≥95%.

682 *2-(2,4-difluorophenyl)-1-(4-(5-fluoropyrimidin-2-yl)piperazin-1-yl)-3-(1H-1,2,4-triazol-*
683 *1-yl)propan-2-ol 15*

684 Light yellow solid (67%); M.P. = 140 – 149 °C; [α]_D26: 0 (c = 0.8 in methanol); ¹H NMR (400
685 MHz, 25 degree, MeOH-d₄) σ: 2.51 (4H, td, J = 4 Hz, 3 Hz), 2.84 (1H, d, J = 14 Hz), 3.02 (dd, J =
686 14 Hz, 2 Hz), 3.65 (4H, t, J = 5 Hz), 4.65 (1H, d, J = 14 Hz), 4.72 (1H, d, J = 14 Hz), 6.87 (1H, tdd,
687 J = 8 Hz, 3 Hz, 1 Hz), 6.94 (1H, ddd, J = 12 Hz, 9 Hz, 3 Hz), 7.50 (1H, td, J = 9 Hz, 7 Hz), 7.76 (1H,
688 s), 8.23 (2H, d, J = 6 Hz), 8.35 (1H, s); ¹³C NMR (101 MHz, 25 degree, MeOH-d₄) σ: 45.5, 55.6,
689 57.5 (d, J = 5 Hz), 64.77 (d, J = 4 Hz), 75.3 (d, J = 6 Hz), 104.9 (dd, J = 28 Hz, 26 Hz), 112.0 (dd,
690 J = 21 Hz, 3 Hz), 127.2 (dd, J = 13 Hz, 4 Hz), 130.9 (dd, J = 10 Hz, 6 Hz), 146.1, 146.3 (d, J = 22
691 Hz), 151.1, 153.2 (d, J = 247 Hz), 160.2 (d, J = 1 Hz), 160.7 (dd, J = 247 Hz, 12 Hz), 164.2 (dd, J
692 = 248 Hz, 12 Hz); HRMS: (ESI+) found 420.1750, ([M+H]⁺) requires 420.17542; Rf : 0.4 (Ethyl
693 acetate); Purity: ≥99%.

694 *2-(2,4-Difluorophenyl)-1-(4-((5-fluoropyrimidin-2-yl)amino)-[1,4'-bipiperidin]-1'-yl)-3-*
695 *(1H-1,2,4-triazol-1-yl)propan-2-ol 16*

696 Yellowish liquid (50%); [α]_D27: 8 (c = 2.5 in methanol); ¹H NMR (400 MHz, 25 degree,
697 methanol-d₄) σ : 1.45 – 1.93 (6H, m), 2.09 – 2.20 (3H, m), 2.35 (1H, td, J = 12 Hz, 2 Hz), 2.55 –
698 2.69 (4H, m), 2.79 (1H, d, J = 14 Hz), 2.90 (1H, m), 3.00 (1H, dd, J = 14 Hz, 1 Hz), 3.14 – 3.20
699 (2H, m), 3.81 (1H, tt, 10 Hz, 4 Hz), 4.60 (1H, d, J = 14 Hz), 4.7 (1H, d, J = 14 Hz), 6.86 (1H, ddd,
700 J = 8 Hz, 8 Hz, 2 Hz), 6.93 (1H, ddd, J = 12 Hz, 9 Hz, 3 Hz), 7.49 (1H, ddd, J = 9 Hz, 9 Hz, 7 Hz),
701 7.76 (1H, s), 8.22 (1H, d, J = 1 Hz), 8.34 (1H, s); ¹³C NMR (101 MHz, 25 degree, MeOH-d₄) σ :
702 28.6 (d, J = 30 Hz), 31.5, 55.0, 55.7, 57.5 (d, J = 5 Hz), 63.6, 64.0 (d, J = 4 Hz), 75.0 (d, J = 5 Hz),
703 79.3, 104.9 (dd, J = 28 Hz, 26 Hz), 112.03 (dd, J = 21 Hz, 3 Hz), 127.3 (dd, J = 12 Hz, 4 Hz), 130.9
704 (dd, J = 9 Hz, 6 Hz), 146.0, 146.7 (d, J = 22 Hz), 151.1, 153.4 (d, J = 246 Hz), 160.4 (d, J = 1 Hz),
705 160.6 (dd, J = 246 Hz, 12 Hz), 164.2 (dd, J = 247 Hz, 12 Hz); m/z: (ESI+) 517.2 ([M+H]⁺); HRMS:
706 (ESI+) found 517.2643, ([M+H]⁺) requires 517.2646; R_f: 0.4 (9 : 1 : 0.1 Ethyl acetate : MeOH :
707 TEA); Purity: \geq 95%.

708 *2-(2,4-difluorophenyl)-1-(8-((5-fluoropyrimidin-2-yl)amino)-2-azaspiro[4.5]decan-2-*
709 *yl)-3-(1H-1,2,4-triazol-1-yl)propan-2-ol 17*

710 Transparent liquid (38%); [α]_D25: 36 (c = 0.6 in methanol); ¹H NMR (400 MHz, 25 degree,
711 MeOH-d₄) σ : 1.11 – 1.25 (1H, m), 1.31 – 1.39 (3H, m), 1.51 (1H, td, J = 7 Hz, 2 Hz), 1.54 – 1.64
712 (3H, m), 1.82 – 1.85 (2H, m), 2.27 (1H, dd, J = 14 Hz, 9 Hz), 2.40 (1H, dd, J = 24 Hz, 9 Hz), 2.51
713 – 2.58 (2H, m), 2.95 (1H, dd, J = 13 Hz, 10 Hz), 3.05 (1H, ddd, J = 23 Hz, 13 Hz, 2 Hz), 3.58 –
714 3.66 (1H, m), 4.59 (1H, dd, J = 14 Hz, 2 Hz), 4.68 (1H, d, J = 14 Hz), 6.86 (1H, td, J = 8 Hz, 2 Hz),
715 6.93 (1H, dddd, J = 11 Hz, 9 Hz, 2 Hz, 2 Hz), 7.46 – 7.53 (1H, m), 7.76 (1H, d, J = 3 Hz), 8.18 (2H,
716 s), 8.35 (1H, s); m/z: (ESI+) 488.2 ([M+H]⁺); HRMS: (ESI+) found 488.2382, ([M+H]⁺) requires
717 488.23802; R_f: 0.3 (Ethyl acetate); Purity: \geq 96%.

718 *2-(2,4-Difluorophenyl)-1-(8-(5-fluoropyrimidin-2-yl)-2,8-diazaspiro[4.5]decan-2-yl)-3-*
719 *(1H-1,2,4-triazol-1-yl)propan-2-ol 18*

720 White solid (55%); M.P. = 110 – 112 °C; [α]_D24: 31 (c = 0.7 in methanol); ¹H NMR (400 MHz,
721 25 degree, methanol-d₄) σ : 1.48 (4H, t, J = 6 Hz), 1.61 (2H, t, J = 7 Hz), 2.37 (1H, d, J = 9 Hz),
722 2.43 (1H, d, J = 9 Hz), 2.62 (2H, t, J = 7 Hz), 2.98 (1H, d, J = 13 Hz), 3.10 (1H, dd, J = 13 Hz, 2 Hz),
723 3.58 – 3.70 (4H, m), 6.86 (1H, tdd, J = 8 Hz, 3 Hz, 1 Hz), 6.93 (1H, ddd, J = 12 Hz, 9 Hz, 3 Hz),
724 7.50 (1H, td, J = 9 Hz, 7 Hz), 7.76 (1H, s), 8.22 (2H, d, J = 1 Hz), 8.35 (1H, s); ¹³C NMR (101 MHz,
725 25 degree, MeOH-d₄) σ : 37.4, 37.9 (d, J = 13 Hz), 41.7, 43.3 (d, J = 2 Hz), 55.8, 57.5 (d, J = 5
726 Hz), 62.8 (d, J = 4 Hz), 67.7, 75.2 (d, J = 6 Hz), 104.8 (dd, J = 28 Hz, 26 Hz), 112.0 (dd, J = 21 Hz,
727 3 Hz), 127.3 (dd, J = 13 Hz, 4 Hz), 131.1 (dd, J = 9 Hz, 6 Hz), 146.1, 146.2 (d, J = 22 Hz), 151.1,
728 152.9 (d, J = 246 Hz), 160.3 (d, J = 1 Hz), 160.8 (dd, J = 246 Hz, 12 Hz), 164.3 (dd, J = 247 Hz, 12
729 Hz); m/z: (ESI+) 474.2 ([M+H]⁺); HRMS: (ESI+) found 474.2220, ([M+H]⁺) requires 474.2224;
730 R_f: 0.2 (Ethyl acetate); Purity: \geq 95%.

731 *2-(2,4-difluorophenyl)-1-(4-((5-fluoropyrimidin-2-yl)amino)azepan-1-yl)-3-(1H-1,2,4-*
732 *triazol-1-yl)propan-2-ol 20*

733 Yellowish liquid (40%); [α]D₂₆: -14 (c = 1.5 in methanol); ¹H NMR (400 MHz, 25 degree,
734 MeOH-d₄) σ : 1.46 – 1.68 (4H, m), 1.81 – 1.89 (2H, m), 2.54 – 2.76 (4H, m), 2.90 (1H, dd, J = 14
735 Hz, 10 Hz), 3.23 (1H, ddd, J = 14 Hz, 3 Hz, 2 Hz), 3.93 (1H, dddd, J = 17 Hz, 13 Hz, 9 Hz, 4 Hz),
736 4.57 (1H, dd, J = 14 Hz, 7 Hz), 4.69 (1H, dd, J = 14 Hz, 1 Hz), 6.84 (1H, td, J = 8 Hz, 3 Hz), 6.89 –
737 6.96 (1H, m), 7.53 (1H, tdd, J = 9 Hz, 11 Hz, 7 Hz), 7.75 (1H, s), 8.19 (2H, s), 8.37 (1H, d, J = 9
738 Hz); ¹³C NMR (101 MHz, 25 degree, MeOH-d₄) σ : 26.1 (d, J = 7 Hz), 34.1 (d, J = 8 Hz), 35.5 (d,
739 J = 35 Hz), 52.3 (d, J = 30 Hz), 54.8 (d, J = 52 Hz), 57.6 (dd, J = 5 Hz, 4 Hz), 58.5 (d, J = 43 Hz),
740 65.3 (dd, J = 30 Hz, 3 Hz), 74.9 (dd, J = 16 Hz, 6 Hz), 104.9 (dd, J = 8 Hz, 2 Hz), 105.1 (d, J = 8
741 Hz), 112.0 (dd, J = 21 Hz, 3 Hz), 127.4 (dd, J = 13 Hz, 4 Hz), 131.2 (dd, J = 9 Hz, 6 Hz), 134.5 (d,
742 J = 64 Hz), 140.2 (d, J = 116 Hz), 146.1 (d, J = 7 Hz), 146.6 (d, J = 22 Hz), 151.1 (d, J = 2 Hz),
743 153.1 (dd, J = 245 Hz, 1 Hz), 160.2 (dd, J = 5 Hz, 1 Hz), 164.2 (dd, J = 249 Hz, 12 Hz); m/z: (ESI+)
744 448.2 ([M+H]⁺); HRMS: (ESI+) found 448.2063, ([M+H]⁺) requires 448.20672; R_f : 0.2 (Ethyl
745 acetate); Purity: \geq 98%.

746 *2-(2,4-Difluorophenyl)-1-(4-(5-fluoropyrimidin-2-yl)-1,4-diazepan-1-yl)-3-(1H-1,2,4-*
747 *triazol-1-yl)propan-2-ol 21*

748 Transparent liquid (52%); [α]D₂₆: 15 (c = 0.7 in methanol); ¹H NMR (400 MHz, 25 degree,
749 methanol-d₄) σ : 1.64 – 1.71(2H, m), 2.54 – 2.69 (2H, m), 2.79 (2H, td, J = 6 Hz, 1 Hz), 2.88 (1H,
750 d, J = 14 Hz), 3.20 (1H, dd, J = 14 Hz, 2 Hz), 3.62 – 3.76 (4H, m), 4.52 (1H, d, J = 14 Hz), 4.64
751 (1H, d, J = 14 Hz), 6.83 (1H, tdd, J = 9 Hz, 3 Hz, 1 Hz), 6.90 (1H, ddd, J = 12 Hz, 9 Hz, 3 Hz), 7.42
752 (1H, td, J = 9 Hz, 7 Hz), 7.75 (1H, s), 8.24 (2H, d, J = 1 Hz), 8.31 (1H, s); ¹³C NMR (101 MHz, 25
753 degree, MeOH-d₄) σ : 27.9, 47.7, 57.4 (d, J = 5 Hz), 57.4, 57.5, 63.3 (d, J = 4 Hz), 75.1 (d, J = 6
754 Hz), 104.8 (dd, J = 28 Hz, 26 Hz), 112.0 (dd, J = 21 Hz, 3 Hz), 127.2 (dd, J = 13 Hz, 4 Hz), 131.0
755 (dd, J = 10 Hz, 6 Hz), 146.0, 146.4 (d, J = 22 Hz), 151.1, 153.0 (d, J = 245 Hz), 159.9 (d, J = 1 Hz),
756 160.6 (dd, J = 246 Hz, 12 Hz), 164.1 (dd, J = 248 Hz, 12 Hz); m/z: (ESI+) 434.1 ([M+H]⁺); HRMS:
757 (ESI+) found 434.1907, ([M+H]⁺) requires 434.1911; R_f : 0.4 (Ethyl acetate); Purity: \geq 96%.

758

759 **Molecular modelling**

760 The protein structures of lanosterol 14-alpha demethylase from *Candida albicans* was
761 obtained from the protein data bank (PDB ID5TZ1) and lanosterol 14-alpha demethylase from
762 *C. auris* (Uniprot ID A0A2H4QC40) was using developed using AlphaFold3. AutoDock SMINA
763 was initially used to identify the preferred binding pockets,⁴⁵ and Pymol was used in parallel
764 to visualise all inhibitors or substrates' binding poses. The parameters were kept at default
765 settings. After the locating most favoured binding sites, GOLD was used for molecular docking
766 of the drug molecules into the selected binding sites of target proteins. GOLD was used for
767 final experiments due to its flexible docking and more reliable and precise binding energy and
768 scoring estimation.

769

770

771

772

773 **MIC susceptibility tests**

774 Microbroth dilution MICs were carried out following EUCAST guidelines. A 96-well plate was
775 filled with 100 μ L RPMI with 2% glucose in each well from column 2 to column 12. 200 μ L of
776 compound diluted down in media from a DMSO stock was then added to the first column of
777 the 96-well plate and diluted two-fold by each column until column 11. Then, 100 μ L of fungi
778 strains from overnight cultures backdiluted to a starting concentration of $\sim 1 \times 10^5$ CFU/mL was
779 added to each well except a blank control row. The plate was incubated at 37 °C for 24 h in
780 the incubator. Fungal growth was measured with a BMG plate reader (FLUOstar Microplate
781 Reader, BMG Labtech) at OD530nm. For azole antifungals, the MIC was defined as the lowest
782 concentration that was able to inhibit $\geq 50\%$ of the drug-free control. DMSO and fluconazole
783 controls were run alongside. All MICs were conducted in triplicate or more until a modal MIC
784 value was obtained, or a range stated if a modal value was not defined.

785

786 **Galleria mellonella survival test**

787 Wax moth larvae (*Galleria mellonella*) were kept on wood chips at 14°C in the dark until use.
788 For experiments, it was assumed that each *Galleria* had a haemolymph volume of 50 μ L and
789 can maximum tolerant 10 μ L of liquid injection. Therefore, the stock solution of each
790 compound was prepared at 6 times of the concentration they were going to be tested. For
791 each compound at each test concentration, 10 μ L of compound stock solution was injected
792 into 10 *Galleria* via the foremost proleg using a Hamilton syringe. Ten of the control larvae
793 were injected with 10 μ L of PSB to control for potential lethal effects from the injection
794 process. After injection, larvae were kept at 37°C inside the petri dishes, and the number of
795 live larvae was recorded every 24 hrs for 5 days. This method was adapted from Wand, et al.⁴⁶

796

797 **Efficacy assay in *G. mellonella* model**

798 *G. mellonella* larvae were injected with 10 μ L of *C. auris* strain TDG1912 at $\sim 1 \times 10^7$ CFU/mL
799 into the first left proleg. Then *G. mellonella* were injected with antimicrobial agent/10% DMSO
800 in PBS in the first right proleg 30 minutes after infection. Controls were injected with PBS
801 alone. 10 larvae were treated per condition, per repeat for a total of 30 larvae per condition
802 across 3 independent repeats. *G. mellonella* were stored at 4 °C, allowed to come to room
803 temperature for at least an hour before the procedure and were used within 2 weeks of the
804 receipt date. *G. mellonella* were assessed for survival every day for 5 days.

805

806 **Time-kill assay**

807 Overnight cultures of fungal strains were back-diluted to a starting concentration of $\sim 1 \times 10^5$
808 CFU/mL into glass universals containing 3 mL of RPMI with 2% glucose media and the drug at
809 the concentration of 4 x MIC₅₀. The glass universals were incubated at 37 °C with shaking at

810 200 rpm for 24 h. Aliquots (20 μ L) were taken out of the glass universals for each tested
811 compound and non-treated (NT) control at six-time points (0, 1, 2, 4, 6 and 24 hours) and
812 Miles-Misra performed to estimate the total number of colony forming units (CFU) per mL.
813 The tests were conducted in triplicate and the average value of log CFU/mL was reported.
814 Along with the synthesised compounds, commercially available antifungal drugs were also
815 tested for comparison. Tested compounds were defined as fungicidal if the loss of fungal
816 population is more than a 3-log reduction in CFU/mL compared to time point 0 h.

817

818 **ASSOCIATED CONTENT**

819 **Supporting information.**

820 Supporting information are available containing:

821 - LC-MS method, NMR spectra and HRMS (PDF)

822 -Molecular formula strings (CSV)

823

824 **AUTHOR INFORMATION**

825 **Corresponding author.**

826 For KMR: e-mail k.miraz.rahman@kcl.ac.uk, phone +44 (0)207 848 1891

827 [For JMS: e-mail mark.sutton@phe.gov.uk](mailto:mark.sutton@phe.gov.uk), phone +44 (0)198 061 2649

828 **ORCID**

829 **Khondaker M. Rahman:** 0000-0001-8566-8648

830 **Notes**

831 The authors declare no competing financial interest.

832

833 **Acknowledgements:**

834 Funding was received from China Schoiarship Council (CSC) and a Medical Research Council
835 Confidence in Concept grant (award code MC_PC_13065).

836

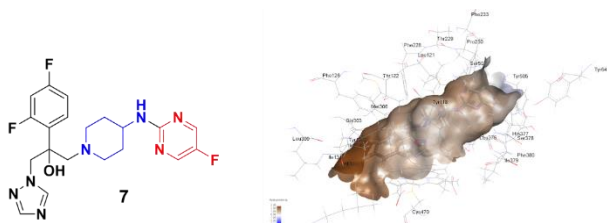
837 **ABBREVIATIONS USED**

838 AcOH, Acetic acid; CDC, Centers for Disease Control and Prevention; calLDM, lanosterol 14 α -
839 demethylase of *C. albicans*; carLDM, lanosterol 14 α -demethylase of *C. auris*; CLSI, Clinical
840 and Laboratory Standards Institute; DCM, Dichloromethane; DIPEA, N, N-

841 Diisopropylethylamine; DMA, Dimethylacetamide; DMSO, Dimethylsulfoxide; DMF, N,N-

842 Dimethylformamide; EtN(Pr-i)₂, N, N-Diisopropylethylamine; ERB, Efflux resistant breaker;
843 EtOH, Ethanol; Equiv., Equivalent; EUCAST, European Committee on Antimicrobial
844 Susceptibility Testing; Et₃N, Triethylamine; ESI, Electrospray Ionization; Fluc, Fluconazole;
845 HCl, Hydrochloric Acid; Hz, Hertz; HPLC, High pressure liquid chromatography; J, Coupling
846 constants; KOtBu, Potassium tert-butoxide; K₂CO₃, Potassium carbonate; LDM, lanosterol
847 14 α -demethylase; LogP, Lipophilicity; LC-MS, Liquid chromatography-mass spectroscopy;
848 MIC, Minimum Inhibitory Concentration; MeOH, Methanol; MeCN, Acetonitrile; MgSO₄,
849 Magnesium sulfate; m/z, mass-to-charge ratio peaks; NaBH(OAc)₃, Sodium
850 triacetoxyborohydride; NaOtBu, Sodium tert-butoxide; NaOH, Sodium hydroxide;
851 Na(OAc)₃BH, Sodium triacetoxyborohydride; NMR, Nuclear magnetic resonance; ORD,
852 optical rotatory dispersion; Pd₂(dba)₃, Tris(dibenzylideneacetone)dipalladium (0); PBS,
853 Phosphate-buffered saline; ppm, parts per million; rac-BINAP, 2,2'-bis(diphenylphosphino)-
854 1,1'-binaphthyl; R_f, retention factors; SAR, Structure Activity Relationship; SNAr, Nucleophilic
855 aromatic substitution; TEA, Triethylamine; TLC, Thin layer chromatography; UV, Ultraviolet;
856 Vori, voriconazole; WHO, World Health Organization; Δ G, Delta G.

857 Graphical abstract



864 **References**

865

- 866 (1) CDC. *Centers for Disease Control and Prevention, National Center for Emerging and*
867 *Zoonotic Infectious Diseases (NCEZID), Division of Foodborne, Waterborne, and*
868 *Environmental Diseases (DFWED), U.S. Department of Health & Human Services.*
869 <https://www.cdc.gov/fungal/antifungal-resistance.html> (accessed 2022 10 November).
- 870 (2) Perlin, D. S.; Rautemaa-Richardson, R.; Alastruey-Izquierdo, A. The global problem of
871 antifungal resistance: prevalence, mechanisms, and management. *Lancet Infect Dis* **2017**, *17*
872 (12), e383-e392. DOI: 10.1016/S1473-3099(17)30316-X From NLM Medline.
- 873 (3) Pappas, P. G.; Rex, J. H.; Sobel, J. D.; Filler, S. G.; Dismukes, W. E.; Walsh, T. J.; Edwards, J.
874 E.; Infectious Diseases Society of, A. Guidelines for treatment of candidiasis. *Clin Infect Dis*
875 **2004**, *38* (2), 161-189. DOI: 10.1086/380796 From NLM Medline.
- 876 (4) Jenks, J. D.; Hoenigl, M. Treatment of Aspergillosis. *J Fungi (Basel)* **2018**, *4* (3). DOI:
877 10.3390/jof4030098 From NLM PubMed-not-MEDLINE.
- 878 (5) Spadari, C. C.; Wirth, F.; Lopes, L. B.; Ishida, K. New Approaches for Cryptococcosis
879 Treatment. *Microorganisms* **2020**, *8* (4). DOI: 10.3390/microorganisms8040613 From NLM
880 PubMed-not-MEDLINE.
- 881 (6) Hendrickson, J. A.; Hu, C.; Aitken, S. L.; Beyda, N. Antifungal Resistance: a Concerning
882 Trend for the Present and Future. *Curr Infect Dis Rep* **2019**, *21* (12), 47. DOI:
883 10.1007/s11908-019-0702-9 From NLM PubMed-not-MEDLINE.
- 884 (7) Stevenson, E. M.; Gaze, W. H.; Gow, N. A. R.; Hart, A.; Schmidt, W.; Usher, J.; Warris, A.;
885 Wilkinson, H.; Murray, A. K. Antifungal Exposure and Resistance Development: Defining
886 Minimal Selective Antifungal Concentrations and Testing Methodologies. *Front Fungal Biol*
887 **2022**, *3*, 918717. DOI: 10.3389/ffunb.2022.918717 From NLM PubMed-not-MEDLINE.
- 888 (8) Morace, G.; Perdoni, F.; Borghi, E. Antifungal drug resistance in *Candida* species. *J Glob*
889 *Antimicrob Resist* **2014**, *2* (4), 254-259. DOI: 10.1016/j.jgar.2014.09.002 From NLM PubMed-
890 not-MEDLINE.
- 891 (9) Whaley, S. G.; Berkow, E. L.; Rybak, J. M.; Nishimoto, A. T.; Barker, K. S.; Rogers, P. D. Azole
892 Antifungal Resistance in *Candida albicans* and Emerging Non-*albicans Candida* Species. *Front*
893 *Microbiol* **2016**, *7*, 2173. DOI: 10.3389/fmicb.2016.02173 From NLM PubMed-not-MEDLINE.
- 894 (10) Goncalves, S. S.; Souza, A. C. R.; Chowdhary, A.; Meis, J. F.; Colombo, A. L. Epidemiology
895 and molecular mechanisms of antifungal resistance in *Candida* and *Aspergillus*. *Mycoses*
896 **2016**, *59* (4), 198-219. DOI: 10.1111/myc.12469 From NLM Medline.
- 897 (11) Chamilos, G.; Kontoyiannis, D. P. Update on antifungal drug resistance mechanisms of
898 *Aspergillus fumigatus*. *Drug Resist Updat* **2005**, *8* (6), 344-358. DOI:
899 10.1016/j.drup.2006.01.001 From NLM Medline.
- 900 (12) Ostrowsky, B.; Greenko, J.; Adams, E.; Quinn, M.; O'Brien, B.; Chaturvedi, V.; Berkow, E.;
901 Vallabhaneni, S.; Forsberg, K.; Chaturvedi, S.; et al. *Candida auris* Isolates Resistant to Three
902 Classes of Antifungal Medications - New York, 2019. *MMWR Morb Mortal Wkly Rep* **2020**, *69*
903 (1), 6-9. DOI: 10.15585/mmwr.mm6901a2 From NLM.
- 904 (13) Chakrabarti, A.; Sood, P. On the emergence, spread and resistance of *Candida auris*:
905 host, pathogen and environmental tipping points. *J Med Microbiol* **2021**, *70* (3). DOI:
906 10.1099/jmm.0.001318 From NLM Medline.
- 907 (14) Osei Sekyere, J. *Candida auris*: A systematic review and meta-analysis of current updates
908 on an emerging multidrug-resistant pathogen. *Microbiologyopen* **2018**, *7* (4), e00578.

909 (15) Cortegiani, A.; Misseri, G.; Fasciana, T.; Giammanco, A.; Giarratano, A.; Chowdhary, A.
910 Epidemiology, clinical characteristics, resistance, and treatment of infections by *Candida*
911 *auris*. *J Intensive Care* **2018**, *6*, 69. DOI: 10.1186/s40560-018-0342-4 From NLM PubMed-
912 not-MEDLINE.

913 (16) Chowdhary, A.; Sharma, C.; Meis, J. F. *Candida auris*: A rapidly emerging cause of
914 hospital-acquired multidrug-resistant fungal infections globally. *Plos Pathog* **2017**, *13* (5),
915 e1006290. DOI: 10.1371/journal.ppat.1006290 From NLM Medline.

916 (17) Rhodes, J.; Fisher, M. C. Global epidemiology of emerging *Candida auris*. *Curr Opin*
917 *Microbiol* **2019**, *52*, 84-89. DOI: 10.1016/j.mib.2019.05.008 From NLM Medline.

918 (18) Lone, S. A.; Ahmad, A. *Candida auris*-the growing menace to global health. *Mycoses*
919 **2019**, *62* (8), 620-637. DOI: 10.1111/myc.12904 From NLM Medline.

920 (19) Mishra, S. K.; Yasir, M.; Willcox, M. *Candida auris*: an emerging
921 antimicrobial-resistant organism with the highest level of concern. *The Lancet Microbe* **2023**,
922 *4* (7), e482-e483. DOI: 10.1016/S2666-5247(23)00114-3 (accessed 2023/11/09).

923 (20) Maphanga, T. G.; Mpembe, R. S.; Naicker, S. D.; Govender, N. P. In Vitro Antifungal
924 Activity of Manogepix and Other Antifungal Agents against South African *Candida auris*
925 Isolates from Bloodstream Infections. *Microbiol Spectr* **2022**, *10* (1), e0171721. DOI:
926 10.1128/spectrum.01717-21 From NLM.

927 (21) Simon, S. P.; Li, R.; Silver, M.; Andrade, J.; Tharian, B.; Fu, L.; Villanueva, D.; Abascal, D.
928 G.; Mayer, A.; Truong, J.; et al. Comparative Outcomes of *Candida auris* Bloodstream
929 Infections: A Multicenter Retrospective Case-Control Study. *Clin Infect Dis* **2023**, *76* (3),
930 e1436-e1443. DOI: 10.1093/cid/ciac735 From NLM Medline.

931 (22) CDC. *Antimicrobial Resistance in Candida*. Centers for Disease Control and Prevention,
932 U.S. Department of Health & Human Services, 2020.
933 <https://www.cdc.gov/fungal/diseases/candidiasis/antifungal-resistant.html#ref-1> (accessed.
934 (23) Toda, M.; Williams, S. R.; Berkow, E. L.; Farley, M. M.; Harrison, L. H.; Bonner, L.;
935 Marceaux, K. M.; Hollick, R.; Zhang, A. Y.; Schaffner, W.; et al. Population-Based Active
936 Surveillance for Culture-Confirmed Candidemia - Four Sites, United States, 2012-2016.
937 *MMWR Surveill Summ* **2019**, *68* (8), 1-15. DOI: 10.15585/mmwr.ss6808a1 From NLM
938 Medline.

939 (24) CDC. *Antibiotic Resistance Threats in the United States*. Centers for Disease Control and
940 Prevention, U.S. Department of Health and Human Services: Atlanta, GA, 2019.

941 (25) Chen, S. C. A.; Sorrell, T. C. Antifungal agents. *Medical Journal of Australia* **2007**, *187* (7),
942 404-409. DOI: <https://doi.org/10.5694/j.1326-5377.2007.tb01313.x>.

943 (26) Wu, J. J.; Pang, K. R.; Huang, D. B.; Tyring, S. K. Therapy of systemic fungal infections.
944 *Dermatol Ther* **2004**, *17* (6), 532-538. DOI: 10.1111/j.1396-0296.2004.04057.x From NLM
945 Medline.

946 (27) Maertens, J. A. History of the development of azole derivatives. *Clin Microbiol Infect*
947 **2004**, *10 Suppl 1*, 1-10. DOI: 10.1111/j.1470-9465.2004.00841.x From NLM Medline.

948 (28) Shafiei, M.; Peyton, L.; Hashemzadeh, M.; Foroumadi, A. History of the development of
949 antifungal azoles: A review on structures, SAR, and mechanism of action. *Bioorganic*
950 *Chemistry* **2020**, *104*, 104240. DOI: <https://doi.org/10.1016/j.bioorg.2020.104240>.

951 (29) Marichal, P.; Bossche, H. Mechanisms of resistance to azole antifungals. *Acta Biochimica*
952 *Polonica* **1995**, *42* (4), 509-516.

953 (30) Lamping, E.; Baret, P. V.; Holmes, A. R.; Monk, B. C.; Goffeau, A.; Cannon, R. D. Fungal
954 PDR transporters: Phylogeny, topology, motifs and function. *Fungal Genetics and Biology*
955 **2010**, *47* (2), 127-142. DOI: <https://doi.org/10.1016/j.fgb.2009.10.007>.

956 (31) Kean, R.; Delaney, C.; Sherry, L.; Borman, A.; Johnson, E. M.; Richardson, M. D.;
957 Rautemaa-Richardson, R.; Williams, C.; Ramage, G. Transcriptome Assembly and Profiling of
958 *Candida auris* Reveals Novel Insights into Biofilm-Mediated Resistance. *mSphere* **2018**, *3* (4).
959 DOI: 10.1128/mSphere.00334-18 From NLM Medline.

960 (32) Tobudic, S.; Kratzer, C.; Presterl, E. Azole-resistant *Candida* spp.—emerging pathogens?
961 *Mycoses* **2012**, *55*, 24-32.

962 (33) Morais Vasconcelos Oliveira, J.; Conceição Oliver, J.; Latércia Tranches Dias, A.; Barbosa
963 Padovan, A. C.; Siqueira Caixeta, E.; Caixeta Franco Ariososa, M. Detection of ERG11
964 Overexpression in *Candida albicans* isolates from environmental sources and clinical isolates
965 treated with inhibitory and subinhibitory concentrations of fluconazole. *Mycoses* **2021**, *64*
966 (2), 220-227. DOI: <https://doi.org/10.1111/myc.13208>.

967 (34) Henry, K. W.; Nickels, J. T.; Edlind, T. D. Upregulation of ERG genes in *Candida* species by
968 azoles and other sterol biosynthesis inhibitors. *Antimicrobial agents and chemotherapy*
969 **2000**, *44* (10), 2693-2700.

970 (35) Nett, J. E.; Crawford, K.; Marchillo, K.; Andes, D. R. Role of Fks1p and Matrix Glucan in
971 *Candida albicans* Biofilm Resistance to an Echinocandin, Pyrimidine, and Polyene.
972 *Antimicrobial Agents and Chemotherapy* **2010**, *54* (8), 3505-3508. DOI:
973 doi:10.1128/aac.00227-10.

974 (36) Bruzual, I.; Riggle, P.; Hadley, S.; Kumamoto, C. A. Biofilm formation by fluconazole-
975 resistant *Candida albicans* strains is inhibited by fluconazole. *Journal of Antimicrobial*
976 *Chemotherapy* **2007**, *59* (3), 441-450. DOI: 10.1093/jac/dkl521 (accessed 11/10/2023).

977 (37) Williamson, B.; Wilk, A.; Guerrero, K. D.; Mikulski, T. D.; Elias, T. N.; Sawh, I.; Cancino-
978 Prado, G.; Gardam, D.; Heath, C. H.; Govender, N. P.; et al. Impact of Erg11 Amino Acid
979 Substitutions Identified in *Candida auris* Clade III Isolates on Triazole Drug Susceptibility.
980 *Antimicrobial Agents and Chemotherapy* **2022**, *66* (1), e01624-01621. DOI:
981 doi:10.1128/AAC.01624-21.

982 (38) Xiang, M. J.; Liu, J. Y.; Ni, P. H.; Wang, S.; Shi, C.; Wei, B.; Ni, Y. X.; Ge, H. L. Erg11
983 mutations associated with azole resistance in clinical isolates of *Candida albicans*. *FEMS*
984 *Yeast Res* **2013**, *13* (4), 386-393. DOI: 10.1111/1567-1364.12042 From NLM Medline.

985 (39) Sagatova, A. A.; Keniya, M. V.; Wilson, R. K.; Monk, B. C.; Tyndall, J. D. Structural insights
986 into binding of the antifungal drug fluconazole to *Saccharomyces cerevisiae* lanosterol 14 α -
987 demethylase. *Antimicrobial agents and chemotherapy* **2015**, *59* (8), 4982-4989.

988 (40) Emami, S.; Ghobadi, E.; Saednia, S.; Hashemi, S. M. Current advances of triazole alcohols
989 derived from fluconazole: Design, in vitro and in silico studies. *European Journal of Medicinal*
990 *Chemistry* **2019**, *170*, 173-194. DOI: <https://doi.org/10.1016/j.ejmech.2019.03.020>.

991 (41) Salehi, F.; Emami, L.; Rezaei, Z.; Khabnadideh, S.; Tajik, B.; Sabet, R. Fluconazole-Like
992 Compounds as Potential Antifungal Agents: QSAR, Molecular Docking, and Molecular
993 Dynamics Simulation. *Journal of Chemistry* **2022**, *2022*, 5031577. DOI:
994 10.1155/2022/5031577.

995 (42) Thamban Chandrika, N.; Shrestha, S. K.; Ngo, H. X.; Tsodikov, O. V.; Howard, K. C.;
996 Garneau-Tsodikova, S. Alkylated Piperazines and Piperazine-Azole Hybrids as Antifungal
997 Agents. *Journal of Medicinal Chemistry* **2018**, *61* (1), 158-173. DOI:
998 10.1021/acs.jmedchem.7b01138.

999 (43) Lebouvier, N.; Giraud, F.; Corbin, T.; Na, Y. M.; Le Baut, G.; Marchand, P.; Le Borgne, M.
1000 Efficient microwave-assisted synthesis of 1-(1H-indol-1-yl)-2-phenyl-3-(1H-1,2,4-triazol-1-
1001 yl)propan-2-ols as antifungal agents. *Tetrahedron Letters* **2006**, *47* (36), 6479-6483. DOI:
1002 <https://doi.org/10.1016/j.tetlet.2006.03.199>.

1003 (44) Singkum, P.; Suwanmanee, S.; Pumeesat, P.; Luplertlop, N. A powerful in vivo alternative
1004 model in scientific research: *Galleria mellonella*. *Acta Microbiol Immunol Hung* **2019**, *66* (1),
1005 31-55. DOI: 10.1556/030.66.2019.001 From NLM Medline.
1006 (45) Koes, D. R.; Baumgartner, M. P.; Camacho, C. J. Lessons Learned in Empirical Scoring
1007 with smina from the CSAR 2011 Benchmarking Exercise. *Journal of Chemical Information and*
1008 *Modeling* **2013**, *53* (8), 1893-1904. DOI: 10.1021/ci300604z.
1009 (46) Wand, M. E.; Muller, C. M.; Titball, R. W.; Michell, S. L. Macrophage and *Galleria*
1010 *mellonella* infection models reflect the virulence of naturally occurring isolates of *B.*
1011 *pseudomallei*, *B. thailandensis* and *B. oklahomensis*. *Bmc Microbiology* **2011**, *11*. DOI:
1012 10.1186/1471-2180-11-11.
1013