

1 **Azole and amphotericin B minimum inhibitory concentrations against *Aspergillus***
2 ***fumigatus*: high agreement between spectrophotometric and visual readings using the**
3 **EUCAST 9.3.2 procedure**

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Abstract

The EUCAST 9.3.2 procedure recommends visual readings of azole and amphotericin B MICs against *Aspergillus* spp. Visual determination of MICs may be challenging. In this work, we aim to obtain and compare visual and spectrophotometric MICs readings of azoles and amphotericin B against *A. fumigatus sensu lato* isolates. Eight hundred and forty-seven *A. fumigatus sensu lato* isolates (*A. fumigatus sensu stricto* [n=828] and cryptic species [n=19]) were tested against amphotericin B, itraconazole, voriconazole, posaconazole, and isavuconazole using the EUCAST EDef 9.3.2 procedure. Isolates were classified as susceptible or resistant/non-wild-type according to the 2020 updated breakpoints. The area of technical uncertainty for the azoles was defined in the updated breakpoints. Visual and spectrophotometric (fungal growth reduction >95% compared to control; read at 540 nm) MICs were compared. Essential (± 1 twofold dilutions) and categorical agreements were calculated. Overall, high essential (97.1%) and categorical (99.6%) agreements were found. We obtained 100% categorical agreements for amphotericin B, itraconazole, and posaconazole and, consequently, no errors were found. Categorical agreements were 98.7% and 99.3% for voriconazole and isavuconazole, respectively. Most of misclassifications for voriconazole and isavuconazole were found to be associated with MIC results falling either in the area of technical uncertainty or in one two-fold dilutions above the breakpoint. Resistance rate was slightly lower when the MICs were obtained by spectrophotometric readings. However, all relevant *cyp51A* mutants were correctly classified as resistant.

58 Spectrophotometric determination of azole and amphotericin B MICs against *A.*
59 *fumigatus sensu lato* isolates may be a convenient alternative to visual endpoint
60 readings.

Introduction

Azoles are the backbone of treatment and prevention of *Aspergillus* spp. diseases and are to date the only available anti-*Aspergillus* oral drugs. The European Society of Clinical Microbiology and Infectious Diseases guidelines recommend itraconazole for the management of patients with chronic pulmonary aspergillosis and allergic bronchopulmonary aspergillosis. Voriconazole and isavuconazole are indicated as the first-line treatment of pulmonary invasive aspergillosis. Voriconazole is also recommended for primary therapy in patients with central nervous system involvement and chronic pulmonary forms of the infection. Posaconazole is recommended for antifungal prophylaxis during prolonged neutropenia in high-risk patients or as salvage therapy in intolerant or non-responding individuals. Finally, liposomal amphotericin B is recommended in settings in which azoles are contraindicated - resistant isolates - and as salvage therapy (1). Some *Aspergillus* species are intrinsically resistant to polyenes (*A. terreus*, *A. nidulans*, and *A. flavus*) or azoles (*A. ustus*) (2). *A. fumigatus sensu lato*, the main etiological agents of aspergillosis, include *A. fumigatus sensu stricto* and cryptic species. Cryptic species commonly show intrinsic resistance to amphotericin B and azoles (3). In contrast, *A. fumigatus sensu stricto* isolates may acquire resistance following exposure to azoles, particularly with environmental azole fungicides (4). Azole resistance in *A. fumigatus sensu stricto* isolates has been increasingly reported worldwide (5-7).

Patients infected by azole-resistant *A. fumigatus sensu lato* isolates show higher mortality than those with azole-susceptible infections (8, 9). Thus, to improve patient care, detection of resistance is of paramount importance. The Clinical and Laboratory Standards Institute (CLSI) and the European Committee on Antimicrobial Susceptibility

85 Testing (EUCAST) proposed standard methods for the study of azole and amphotericin B
86 susceptibility of *Aspergillus* spp isolates. The EUCAST EDef 9.3.2 procedure includes
87 clinical breakpoints to classify isolates either as susceptible or resistant and
88 recommends visual determination of MICs (10). Visual inspection may be challenging
89 and spectrophotometric readings may facilitate MIC determination and overcome
90 subjectivity. However, there is a limited number of studies using the EUCAST
91 methodology in which azole MICs against *A. fumigatus sensu lato* obtained by visual and
92 spectrophotometric readings are compared; furthermore, the studies are thwarted by a
93 low number of isolates and antifungal drugs tested (11-14).

94 We recently conducted a Spanish multicenter study of azole-resistance in which
95 847 *A. fumigatus sensu lato* clinical isolates were collected between February 15 and
96 May 14, 2019 (15). Taking advantage of the large number of isolates, the objective in
97 this work is to report and compare azole and amphotericin B MICs using visual and
98 spectrophotometric readings following the EUCAST EDef 9.3.2 procedure.

99

100 Results

101 Tables 2 to 6 show MIC distributions of amphotericin B, itraconazole,
102 posaconazole, voriconazole, and isavuconazole against the 847 isolates by
103 regular/stringent visual and spectrophotometric readings. MICs against QC strains were
104 within the acceptable limit.

105 Agreement between MICs by regular visual and spectrophotometric readings.

106 Overall, both MIC endpoints showed high essential (97.1%) and categorical agreements
107 (99.6%). Essential agreements for individual drugs were as follows: amphotericin B

108 98.8%, itraconazole 94.8%, posaconazole 97.3%, voriconazole 98.3%, and isavuconazole
109 96.1% (Table 7).

110 Categorical agreements for amphotericin B, itraconazole, and posaconazole were
111 100% and, consequently, resistance rates for both MIC endpoints were identical.
112 Categorical agreement for voriconazole was 98.7% and the rate of resistance was
113 slightly lower when spectrophotometric readings were used for MIC determination.
114 Very major errors (n=6, 0.7%) and major errors (n=3, 0.4%) for voriconazole occurred in
115 *A. fumigatus sensu stricto* isolates with MIC results falling in the ATU (MIC=2 mg/L). In
116 cryptic species, very major errors occurred in two *N. udagawae* isolates (10.5%), one of
117 them with MIC results falling in the ATU. Categorical agreement for isavuconazole was
118 99.3% and the rate of resistance was slightly lower when spectrophotometric readings
119 were used for MIC determination. Very major errors in isavuconazole occurred in three
120 *A. fumigatus sensu stricto* isolates and in three cryptic species isolates (two *N.*
121 *udagawae* and one *A. fumigatiaffinis*). With the exception of the *A. fumigatiaffinis*
122 isolate, very major errors for isavuconazole (n=5) were detected in isolates with MIC
123 results in the ATU, which also revealed very major errors for voriconazole (Table 1S).
124 None of the six isolates for which very major errors were detected in the azole
125 categorical classification harboured relevant *cyp51A* mutations (Table 1S).

126 **Agreement between MICs obtained by regular/stringent visual readings.** Overall,
127 both visual MIC endpoints showed high essential (97.7%) and categorical agreements
128 (96.7%). Essential agreements for individual drugs were above 98% (itraconazole
129 [98.9%], posaconazole [98.7%], and isavuconazole [98.6%]) with the exception of
130 voriconazole (94.4%) (Table 7).

131 Categorical agreements for itraconazole and posaconazole were 99.4% (Table 7).
132 Resistance rates obtained by both MIC endpoints were identical in *A. fumigatus sensu*
133 *stricto*, but slightly higher with stringent visual readings in cryptic species. This led to
134 major errors for both drugs in five isolates (three *A. lentulus*, one *A. novofumigatus* and
135 one *A. fumigatiaffinis*). Although posaconazole MICs by both visual readings were
136 identical (MIC=0.25 mg/L, ATU), the categorical classification differed due to the MICs of
137 itraconazole in four out of the five isolates (Table 2S). Percentage of voriconazole
138 resistance was overestimated with the visual stringent endpoint (6.6% vs 15.8%).
139 Categorical agreement was 90.8%. Major errors were found exclusively in *A. fumigatus*
140 *sensu stricto* isolates (n=78), in MIC results falling in the ATU. Likewise, the rate of
141 isavuconazole resistance was overestimated when visual stringent endpoint was used,
142 although to a lesser extent than in the case of voriconazole (4.1% vs 4.4%). Categorical
143 agreement was 97.2%. Major errors were found in *A. fumigatus sensu stricto* isolates
144 (n=22) and in two isolates of cryptic species (*N. tsurutae* and *A. fumigatiaffinis*; Table
145 2S). Similarly, most misclassifications (23/24 isolates) were associated to MIC results
146 falling in the ATU and mostly affected isolates in which major errors for voriconazole
147 were detected (21/24 isolates). Since stringent visual readings shifted azole MICs to
148 higher values, no very major errors were found.

149

150 Discussion

151 In this study we show that MICs of azoles and amphotericin B against *A.*
152 *fumigatus* obtained either by spectrophotometric or regular visual readings have very
153 high essential and categorical agreement.

154 The increase of resistant *A. fumigatus* isolates worldwide has promoted antifungal
155 susceptibility testing (5). Azole resistance in *A. fumigatus* may occur during azole
156 therapy or exposure to azole fungicides in the environment (4). Furthermore, cryptic
157 species commonly show intrinsic resistance to amphotericin B and azoles (3). Although
158 the EUCAST EDef 9.3.2 procedure recommends visual inspection for azole and
159 amphotericin B MIC setting against *Aspergillus* species, spectrophotometric readings
160 may offer objectivity, quick automated readings, and overall better performance.
161 Previous studies comparing spectrophotometric and visual readings showed excellent
162 essential (92%-97%) and categorical (93-99%) agreements (11-14). Some of the studies
163 used the CLSI methodology and were undermined by the limited number of *A.*
164 *fumigatus sensu stricto* tested isolates (up to 133 isolates), the absence of both cryptic
165 species isolates and *cyp51A* mutants, and a low number of studied antifungal
166 drugs(amphotericin B and itraconazole) (12-14). One of the studies, in which the EUCAST
167 method was used, included the four anti-mold triazoles (itraconazole, posaconazole,
168 voriconazole, and isavuconazole) and a low number of *A. fumigatus sensu stricto*
169 isolates (n= 88). The work did not assess cryptic species, although 15 isolates with
170 *cyp51A* mutations including isolates with the dominant substitutions TR₃₄/L98H, G54,
171 M220, among others, were examined. Furthermore, since EUCAST has recently changed
172 azole breakpoints against *Aspergillus fumigatus sensu lato*, a validation of
173 spectrophotometric readings including a large number of isolates classified according to
174 the updated EUCAST breakpoints is needed.

175 We recently conducted a survey of azole resistance in *A. fumigatus sensu lato*
176 isolates collected in Spain in 2019 (15). Taking advantage of the large number of isolates
177 collected (n=847), we obtained and compared MICs using visual and spectrophotometric

178 readings. Nineteen strains were identified as cryptic species and 45 *A. fumigatus sensu*
179 *stricto* proved to be azole resistant, being TR₃₄-L98H the dominant mechanism of
180 resistance. Both MIC endpoints show high essential/categorical agreements for
181 amphotericin B (98.8/100), itraconazole (94.8%/100%), posaconazole (97.3%/100%),
182 voriconazole (98.3%/98.7%), and isavuconazole (96.1%/99.3%). No errors were found in
183 amphotericin B, itraconazole, and posaconazole. Most misclassifications for
184 voriconazole and isavuconazole are linked with MIC results falling either in ATU (10/12
185 isolates) or in just one two-fold dilutions above the breakpoint (2/12 isolates; MIC = 4
186 mg/L). Cross-resistance between voriconazole and isavuconazole is the norm in *A.*
187 *fumigatus sensu stricto* (16). Using voriconazole as a surrogate marker,
188 spectrophotometric readings resulted in misdetection of voriconazole resistance in six *A.*
189 *fumigatus sensu stricto* isolates with either a wild-type *cyp51A* gene or genetic
190 polymorphisms of dubious clinical implications (Table 1S).

191 The EUCAST has recently reviewed the antifungal breakpoints against *A.*
192 *fumigatus sensu lato*. Breakpoints for amphotericin B, itraconazole, voriconazole, and
193 posaconazole were lowered, while for isavuconazole it was increased (17). Based on the
194 updated breakpoints, spectrophotometric MIC readings led to correctly classify all
195 isolates with relevant *cyp51A* mutations as resistant. Interpretation uncertainties
196 regarding MIC values may arise in the ATU, a newly introduced term, when the
197 breakpoint of wild-type isolates and mutant isolates converge (17). Isolates with
198 posaconazole and isavuconazole MICs of 0.25 mg/L and 2 mg/L, respectively, cannot be
199 automatically reported as susceptible or resistant. MIC determinations using
200 spectrophotometric readings frequently led to the underestimation of resistance for
201 MIC values falling in the ATU. Here, we were able to easily clarify misclassifications by

202 visually inspecting the tray. False resistance was detected in four *A. fumigatus sensu*
203 *stricto* isolates for which a voriconazole MIC of 2 mg/L was determined by
204 spectrophotometric readings.

205 Higher mortality rate is observed in patients infected with azole-resistant *A.*
206 *fumigatus sensu lato* isolates. Resistance is frequently caused by mutations in the
207 *cyp51A* gene, some of which are associated with a pan-triazole-resistant phenotype
208 (high-level resistance) (8). Some phenotypes only affect the activity of a single azole, or
209 several triazoles with similar molecular structure, and the MIC is close to the clinical
210 breakpoint, resulting in low-level resistance (18). Previous studies have shown that
211 patients infected with low-level voriconazole-resistant *A. fumigatus* (MIC=2 mg) and
212 low-level isavuconazole-resistant *A. fumigatus* (MIC=2 mg) may be treated with
213 voriconazole or isavuconazole, respectively, provided that higher doses are
214 administered (19, 20). In cryptic species, very major errors in voriconazole and
215 isavuconazole were detected (Table 1S).

216 Visual MIC readings may be challenging and taking small colonies into account
217 (stringent visual readings) may result in overestimation of resistance rates and increase
218 the MIC of the isolates one or two two-fold dilutions, particularly for voriconazole. Thus,
219 major errors in voriconazole and isavuconazole (MIC results falling in the ATU) against *A.*
220 *fumigatus sensu stricto* may be detected. Correct classification of relevant *cyp51A* gene
221 mutants was achieved by stringent visual readings.

222 We conclude that spectrophotometric determination is a useful alternative to
223 visual inspection of azole and amphotericin MICs against *A. fumigatus sensu stricto*.
224 Both endpoints show high essential and categorical agreements. Future studies

225 including more isolates from cryptic species and *A. fumigatus sensu stricto* with other
226 kind of *cyp51A* mutations are warranted.

227

228 **Materials and methods**

229 **Samples.** Eight hundred and forty-seven *A. fumigatus sensu lato* clinical isolates,
230 identified by MALDI-TOF, were collected in a 30-hospital survey conducted in Spain (15).
231 Azole-resistant isolates (n=45 *A. fumigatus sensu stricto* and n=19 cryptic species) were
232 molecularly identified. Isolate distribution as per species identification was as follows: *A.*
233 *fumigatus sensu stricto* (n=828), *A. lentulus* (n=6), *A. fumigatiaffinis* (n=5), *Neosartorya*
234 *tsurutae* (n=3), *N. udagawae* (n=2), *A. novofumigatus* (n=2), and *A. thermomutatus*
235 (n=1).

236 The *cyp51A* gene sequence from 45 azole-resistant *A. fumigatus sensu stricto*
237 isolates carried the following mutations: TR₃₄-L98H (n=24), G54R (n=5),
238 TR₄₆/Y121F/T289A (n=1), F46Y/M172V/N248T/D255E/E427K (n=2),
239 F46Y/M172V/N248T/D255E/E416Q/E427K (n=1), F165L (n=1), S496L (n=1), and wild-
240 type *cyp51A* gene (n=10).

241 **EUCAST Antifungal susceptibility testing.** All isolates were subcultured on potato
242 dextrose agar or Sabouraud dextrose agar and incubated at 35°C for 2 to 5 days. Isolates
243 from cryptic species were incubated long enough to assure filtered conidia suspensions
244 reaching a sufficient inoculum (equivalent to McFarland 0.5 using a spectrophotometer).
245 Isolate antifungal susceptibilities to amphotericin B, itraconazole, voriconazole,
246 posaconazole, and isavuconazole were determined following the EUCAST EDef 9.3.2
247 procedure (21). The inoculated trays were incubated for 48 hours at 35 °C and MICs

248 obtained using a visual endpoint (defined as the concentration that completely inhibits
249 fungal growth) and a spectrophotometric endpoint ($\geq 95\%$ inhibition of fungal growth
250 compared to the drug-free control and read at 540 nm, as described elsewhere) (11).
251 Although the EUCAST EDef 9.3.2 procedure recommends ignoring single colonies on the
252 surface, sometimes it is difficult to discern real growth from small colonies. Thus, we
253 interpreted visual MICs using two endpoints: regular endpoint (very tiny growth was
254 disregarded) or stringent endpoint (a totally clear well), as exemplified in Figure 1.
255 Quality control (QC) was ensured by testing the *A. flavus* ATCC 204304 and *A. fumigatus*
256 ATCC 204305 strains (amphotericin B, itraconazole, voriconazole, and posaconazole),
257 and *Candida krusei* ATCC 6258 and *C. parapsilosis* ATCC 22019 (isavuconazole).

258 **Data analysis.** Regular visual endpoint MICs were assumed as the gold standards
259 and compared against MICs obtained by other endpoints; MICs (percentage) within ± 1
260 two-fold dilutions were considered to be in essential agreement. Isolates were classified
261 as resistant/non-wild-type according to the updated 2020 EUCAST breakpoints (Table 1);
262 intermediate category for amphotericin B and azoles and the category of “susceptible
263 increased exposure” are no longer available, and the term area of technical uncertainty
264 (ATU) for the four azoles has been defined (10). ATU is a warning to laboratories on an
265 uncertainty needing attention before reporting the results and represents an area of
266 confluence of both wild-type and mutant isolates particularly for voriconazole,
267 posaconazole, and isavuconazole. MIC results in the ATU were interpreted as follows:
268 itraconazole and voriconazole (always resistant), posaconazole (resistant only if the
269 isolate was also resistant to itraconazole), isavuconazole (resistant only if the isolate was
270 also resistant to voriconazole). Categorical agreement between the three endpoints was
271 assessed. The endpoints were in categorical agreement when the results were in the

272 same susceptibility category (regardless of the MIC). Errors were defined as very major
273 errors (false susceptibility) when the gold standard endpoint classified an isolate as
274 resistant and the other endpoints as susceptible, and as major errors (false resistance)
275 when the gold standard endpoint classified an isolate as susceptible and the other
276 endpoints as resistant (22).

277

278

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417 **Tables and figures**418 **Table 1.** Azole and amphotericin B breakpoints chosen to classify *Aspergillus fumigatus*419 *sensu lato* isolates as susceptible, resistant, or non-wild-type (17)

Drug	ECOFF (mg/L)	Clinical breakpoints (mg/L)		
	WT ≤	S ≤	R ≥	ATU
Amphotericin B	1	1	2	ND
Itraconazole	1	1	2	2
Posaconazole	0.25	0.125	0.25	0.25
Voriconazole	1	1	2	2
Isavuconazole	2	1	4	2

420

421 **ECOFF**, epidemiological cut-off value; **WT**, wild-type; **S**, susceptible; **R**, resistant; **ATU**, area of technical
422 uncertainty; **ND**, not defined.423 Isolates with itraconazole and voriconazole MICs results that fall in the ATU were always considered
424 resistant; isolates with isavuconazole MICs and posaconazole MICs results that fall in the ATU were
425 considered as resistant when voriconazole-resistant or itraconazole-resistant, respectively.

426 **Table 2.** MIC distributions of amphotericin B against the 847 *A. fumigatus sensu lato* isolates. MIC distributions by regular visual readings and
427 their correspondent rates of resistance are reported elsewhere (15)

then, corresponding rates of resistance are reported elsewhere [15]

428

Amphotericin B	MIC distributions (number of isolates for each MIC, in mg/L)												No. of resistant isolates (%) ^a
	0.008	0.016	0.03	0.06	0.125	0.25	0.5	1	2	4	8	≥ 16	
<i>A. fumigatus sensu lato</i> (n=847)													
Regular visual readings	0	0	1	5	70	407	306	45	<u>7</u>	<u>5</u>	<u>1</u>	<u>0</u>	13 (1.5)
Spectrophotometric readings	0	0	0	7	96	452	248	31	<u>6</u>	<u>5</u>	<u>0</u>	<u>2</u>	13 (1.5)
<i>A. fumigatus sensu stricto</i> (n=828)													
Regular visual readings	0	0	1	5	68	407	305	42	<u>0</u>	<u>0</u>	<u>0</u>	<u>0</u>	0 (0) 432
Spectrophotometric readings	0	0	0	7	94	452	246	29	<u>0</u>	<u>0</u>	<u>0</u>	<u>0</u>	0 (0)
Cryptic species (n=19)													
Regular visual readings	0	0	0	0	2	0	1	3	<u>7</u>	<u>5</u>	<u>1</u>	<u>0</u>	13 (68.4)
Spectrophotometric readings	0	0	0	0	2	0	2	2	<u>6</u>	<u>5</u>	<u>0</u>	<u>2</u>	13 (68.4)
Isolates with tandem repeats (n=25)													
Regular visual readings	0	0	0	0	2	10	13	0	<u>0</u>	<u>0</u>	<u>0</u>	<u>0</u>	0 (0)
Spectrophotometric readings	0	0	0	0	3	10	12	0	<u>0</u>	<u>0</u>	<u>0</u>	<u>0</u>	0 (0)

437 Underlined values indicate non-wild-type isolates according to tentative ECOFFs and values in bold indicate resistant isolates (EUCAST Breakpoint table v 10.0, 2020).

438 ^{*}Identical number of resistant isolates and non-wild-type isolates were obtained

439

440 **Table 3.** MIC distributions of itraconazole against the 847 *A. fumigatus sensu lato* isolates. MIC distributions by regular visual
 441 readings and their correspondent rates of resistance are reported elsewhere (15)

Itraconazole	MIC distributions (number of isolates for each MIC, in mg/L)												No. of resistant isolates (%) *
	0.008	0.016	0.03	0.06	0.125	0.25	0.5	1	2	4	8	≥ 16	
<i>A. fumigatus sensu lato</i> (n=847)													444
Regular visual readings	0	0	0	0	26	427	328	21	<u>2</u>	<u>2</u>	<u>2</u>	<u>39</u>	45 (5.3)
Stringent visual readings	0	0	0	0	10	269	459	59	<u>2</u>	<u>2</u>	<u>2</u>	<u>44</u>	50 (5.9)
Spectrophotometric readings	0	0	15	22	49	412	287	17	<u>3</u>	<u>1</u>	<u>1</u>	<u>40</u>	45 (5.3)
<i>A. fumigatus sensu stricto</i> (n=828)													447
Regular visual readings	0	0	0	0	26	426	326	15	<u>2</u>	<u>1</u>	<u>1</u>	<u>31</u>	35 (4.2)
Stringent visual readings	0	0	0	0	10	269	457	57	<u>2</u>	<u>1</u>	<u>1</u>	<u>31</u>	35 (4.2)
Spectrophotometric readings	0	0	15	22	49	410	282	15	<u>3</u>	<u>0</u>	<u>1</u>	<u>31</u>	35 (4.2)
Cryptic species (n=19)													449
Regular visual readings	0	0	0	0	0	1	2	6	<u>0</u>	<u>1</u>	<u>1</u>	<u>8</u>	10 (52.6)
Stringent visual readings	0	0	0	0	0	0	2	2	<u>0</u>	<u>1</u>	<u>1</u>	<u>13</u>	15 (78.9)
Spectrophotometric readings	0	0	0	0	0	2	5	2	<u>0</u>	<u>1</u>	<u>0</u>	<u>9</u>	10 (52.6)
Isolates with tandem repeats (n=25)													454
Regular visual readings	0	0	0	0	0	0	1	0	<u>0</u>	<u>0</u>	<u>0</u>	<u>24</u>	24 (96)
Stringent visual readings	0	0	0	0	0	0	0	1	<u>0</u>	<u>0</u>	<u>0</u>	<u>24</u>	24 (96)
Spectrophotometric readings	0	0	0	0	0	0	1	0	<u>0</u>	<u>0</u>	<u>0</u>	<u>24</u>	24 (96)

455 Values shaded in grey indicate MICs in the area of technical uncertainty (ATU) and were classified as resistant isolates. Underlined values indicate non-wild-type isolates
 456 according to tentative ECOFFs and values in bold indicate resistant isolates (EUCAST Breakpoint table v 10.0, 2020).

457 *Identical number of resistant isolates and non-wild-type isolates were obtained.

458

459 **Table 4.** MIC distributions of posaconazole against the 847 *A. fumigatus sensu lato* isolates. MIC distributions by regular visual readings and
 460 their correspondent rates of resistance or non-wild-type isolates are reported elsewhere (15)

Posaconazole	MIC distributions (number of isolates for each MIC, in mg/L)												No. of isolates (%)	
	0.008	0.016	0.03	0.06	0.125	0.25	0.5	1	2	4	8	≥ 16	Resistant	Non-wild-type
<i>A. fumigatus sensu lato</i> (n=847)														
Regular visual readings	0	2	46	441	279	42	27	3	0	1	0	6	46 (5.4)	37 (4.4)
Stringent visual readings	0	1	18	268	399	119	23	12	0	1	0	6	51 (6)	42 (5)
Spectrophotometric readings	1	1	70	476	225	36	28	4	1	1	0	4	47 (5.5)	38 (4.5)
<i>A. fumigatus sensu stricto</i> (n=828)														
Regular visual readings	0	2	46	440	278	32	20	3	0	1	0	6	34 (4.1)	30 (3.6)
Stringent visual readings	0	1	18	268	398	111	13	12	0	1	0	6	34 (4.1)	32 (3.9)
Spectrophotometric readings	1	1	70	475	222	27	22	4	1	1	0	4	34 (4.1)	32 (3.9)
Cryptic species (n=19)														
Regular visual readings	0	0	0	1	1	10	7	0	0	0	0	0	12 (63.1)	7 (36.8)
Stringent visual readings	0	0	0	0	1	8	10	0	0	0	0	0	17 (89.5)	10 (52.6)
Spectrophotometric readings	0	0	0	1	3	9	6	0	0	0	0	0	12 (63.2)	6 (31.6)
Isolates with tandem repeats (n=25)														
Regular visual readings	0	0	0	0	0	3	18	2	0	0	0	2	24 (96)	22 (88)
Stringent visual readings	0	0	0	0	0	1	11	11	0	0	0	2	24 (96)	24 (96)
Spectrophotometric readings	0	0	0	0	0	1	19	4	1	0	0	0	24 (96)	24 (96)

461 Values shaded in grey indicate MICs in the area of technical uncertainty (ATU). MIC results against *A. fumigatus sensu lato* falling in the ATU were translated to resistant as
 462 follows: regular visual readings (n=9/42), stringent visual readings (n=9/119), and spectrophotometric readings (n=9/36). Underlined values indicate non-wild-type isolates
 463 according to tentative ECOFFs and values in bold indicate resistant isolates (EUCAST Breakpoint table v 10.0, 2020).
 464
 465

466 **Table 5.** MIC distributions of voriconazole against the 847 *A. fumigatus sensu lato* isolates. MIC distributions by regular visual
 467 readings and their correspondent rates of resistance are reported elsewhere (15)

Voriconazole	MIC distributions (number of isolates for each MIC, in mg/L)												No. of resistant isolates (%) *
	0.008	0.016	0.03	0.06	0.125	0.25	0.5	1	2	4	8	≥ 16	
<i>A. fumigatus sensu lato</i> (n=847)													
Visual reading	0	0	0	0	3	82	529	177	<u>19</u>	<u>27</u>	<u>7</u>	<u>3</u>	56 (6.6)
Stringent visual readings	0	0	0	0	1	17	302	394	<u>87</u>	<u>31</u>	<u>11</u>	<u>4</u>	133 (15.7)
Spectrophotometric readings	0	0	0	0	4	138	500	154	<u>18</u>	<u>24</u>	<u>6</u>	<u>3</u>	51 (6)
<i>A. fumigatus sensu stricto</i> (n=828)													
Visual reading	0	0	0	0	3	82	529	176	<u>13</u>	<u>19</u>	<u>3</u>	<u>3</u>	38 (4.6) 474
Stringent visual readings	0	0	0	0	1	17	302	393	<u>86</u>	<u>18</u>	<u>7</u>	<u>4</u>	115 (13.9)
Spectrophotometric readings	0	0	0	0	4	138	498	153	<u>11</u>	<u>18</u>	<u>4</u>	<u>2</u>	35 (4.2) 475
Cryptic species (n=19)													
Visual reading	0	0	0	0	0	0	0	1	<u>6</u>	<u>8</u>	<u>4</u>	<u>0</u>	18 (94.7)
Stringent visual readings	0	0	0	0	0	0	0	1	<u>1</u>	<u>13</u>	<u>4</u>	<u>0</u>	18 (94.7)
Spectrophotometric readings	0	0	0	0	0	0	2	1	<u>7</u>	<u>6</u>	<u>2</u>	<u>1</u>	16 (84.2)
Isolates with tandem repeats (n=25)													
Visual reading	0	0	0	0	0	0	0	0	<u>5</u>	<u>15</u>	<u>2</u>	<u>3</u>	25 (100)
Stringent visual readings	0	0	0	0	0	0	0	0	<u>1</u>	<u>14</u>	<u>7</u>	<u>3</u>	25 (100)
Spectrophotometric readings	0	0	0	0	0	0	0	0	<u>6</u>	<u>14</u>	<u>3</u>	<u>2</u>	25 (100)

482 Values shaded in grey indicate MICs in the area of technical uncertainty (ATU) and were classified as resistant isolates. Underlined values indicate non-wild-type isolates
 483 according to tentative ECOFFs and values in bold indicate resistant isolates (EUCAST Breakpoints table v 10.0, 2020).

484 *The number of resistant isolates and non-wild-type isolates were identical.

485

Table 6. MIC distributions of isavuconazole against the 847 *A. fumigatus sensu lato* isolates. MIC distributions by regular visual readings and their correspondent rates of resistance/non wild-type isolates were reported elsewhere (15)

Isavuconazole	MIC distributions (number of isolates for each MIC, in mg/L)												No. of isolates (%)	
	0.008	0.016	0.03	0.06	0.125	0.25	0.5	1	2	4	8	≥ 16	Resistant	Non-wild-type
<i>A. fumigatus sensu lato</i> (n=847)														
Regular visual readings	0	0	0	0	0	13	440	333	26	14	17	4	48 (5.6)	35 (4.1)
Stringent visual readings	0	0	0	0	0	2	146	572	90	11	21	5	72 (8.5)	37 (4.4)
Spectrophotometric readings	0	0	0	17	4	14	434	314	31	12	18	3	42 (5)	33 (3.9)
<i>A. fumigatus sensu stricto</i> (n=828)														
Regular visual readings	0	0	0	0	0	13	440	327	18	10	16	4	35 (4.2)	30 (3.6)
Stringent visual readings	0	0	0	0	0	2	146	568	80	7	20	5	57 (6.9)	32 (3.9)
Spectrophotometric readings	0	0	0	17	4	14	433	306	24	9	18	3	32(3.9)	30 (3.6)
Cryptic species (n=19)														
Regular visual readings	0	0	0	0	0	0	0	6	8	4	1	0	13 (68.4)	5 (26.3)
Stringent visual readings	0	0	0	0	0	0	0	4	10	4	1	0	15 (78.9)	5 (26.3)
Spectrophotometric readings	0	0	0	0	0	0	1	8	7	3	0	0	10 (52.6)	3 (15.8)
Isolates with tandem repeats (n=25)														
Regular visual readings	0	0	0	0	0	0	0	0	1	6	14	4	25 (100)	24 (96)
Stringent visual readings	0	0	0	0	0	0	0	0	0	2	18	5	25 (100)	25 (100)
Spectrophotometric readings	0	0	0	0	0	0	0	0	0	7	15	3	25 (100)	25 (100)

Values shaded in grey indicate MICs in the area of technical uncertainty (ATU). MIC results against *A. fumigatus sensu lato* falling in the ATU were translated to resistant as follows: regular visual readings (n=13/26), stringent visual readings (n=35/90), and spectrophotometric readings (n=9/31). Underlined values indicate non-wild-type isolates according to tentative ECOFFs and values in bold indicate resistant isolates (EUCAST Breakpoint table v 10.0, 2020)

506 **Table 7.** Essential and categorical agreement between MICs by visual (regular and stringent) and spectrophotometric readings

MIC readings	<i>A. fumigatus sensu lato</i> (%)				<i>A. fumigatus sensu stricto</i> (%)				Cryptic species (%)			
	Essential agreement*	Categorical agreement*	VME	ME	Essential agreement	Categorical agreement	VME	ME	Essential agreement	Categorical agreement	VME	ME
Amphotericin B												
Regular visual vs spectrophot. readings	98.8	100	0	0	99	100	0	0	89.5	100	0	0
Itraconazole												
Regular visual vs spectrophot. readings	94.8	100	0	0	95.1	100	0	0	78.9	100	0	0
Regular visual vs stringent readings	98.9	99.4	0	0.6	99.6	100	0	0	68.4	73.7	0	26.3
Posaconazole												
Regular visual vs spectrophot. readings	97.3	100	0	0	97.1	100	0	0	100	100	0	0
Regular visual vs stringent readings	98.7	99.4	0	0.6	98.7	100	0	0	100	73.7	0	26.3
Voriconazole												
Regular visual vs spectrophot. readings	98.3	98.7	0.9	0.3	98.4	98.9	0.7	0.4	94.7	89.5	10.5	0
Regular visual vs stringent readings	94.4	90.8	0	9.2	94.3	90.6	0	9.4	100	100	0	0
Isavuconazole												
Regular visual vs spectrophot. readings	96.1	99.3	0.7	0	96.3	99.6	0.4	0	89.5	84.2	15.8	0
Regular visual vs stringent readings	98.6	97.2	0	2.8	98.5	97.3	0	2.7	100	89.5	0	10.5

507 *Regular visual endpoint MICs were assumed as the gold standards and compared against MICs obtained by other endpoints; MICs (percentage) within ± 1 two-fold
508 dilutions were considered to be in essential agreement. Isolates were classified as resistant/non-wild-type according to the updated 2020 EUCAST breakpoints. The
509 endpoints were in categorical agreement when the results were in the same susceptibility category (regardless of the MIC). VME: very major error (false susceptibility); ME:
510 major error (false resistance).
511
512

Figure 1. Example of fungal growth of an *A. fumigatus sensu stricto* isolate in the presence of itraconazole (1a), posaconazole (1b), voriconazole (1c), and isavuconazole (1d). Two MIC endpoints were used: the regular endpoint (gold standard) where tiny small colonies were disregarded (wells surrounded by rings of dashed lines) and the stringent endpoint where the tiny colonies were taken into account (wells surrounded by rings of solid lines)

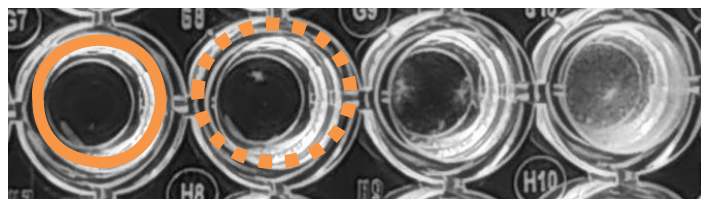
1a

0.5 mg/L 0.25 mg/L 0.125 mg/L



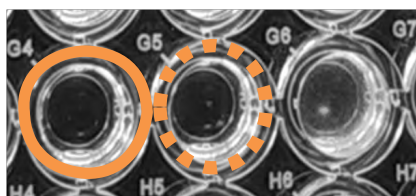
1b

0.125 mg/L 0.06 mg/L 0.03 mg/L 0.015 mg/L



1c

1 mg/L 0.5 mg/L 0.25 mg/L



1d

2 mg/L 1 mg/L 0.5 mg/L 0.25 mg/L 0.125 mg/L

