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# Relation of the polymorphism of *cyp*51A sequence and the susceptibility of *Aspergillus fumigatus* isolates to triazoles determined by commercial gradient test (Etest) and by reference methods

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The aim of this study was to evaluate the accuracy of commercial gradient test (Etest) in the detection of triazole resistant Aspergillus fumigatus isolates using reference microdilution methods and the analysis of sequences of the cyp 51A gene. The study was performed on twenty clinical isolates which were identified as Aspergillus fumigatus based on the DNA sequences of the ITS1-2 fragment of ribosomal DNA and the β-tubulin gene, out of them seventeen isolates showed wild-type cvp51A sequence and three were positive for the mutation TR34/L98H. All isolates were tested for the susceptibility to itraconazole (ITZ), voriconazole (VOR) and posaconasole (POS) using microdilution methods, according to EUCAST and CLSI protocols, as well as using Etest. The results of microdilution and Etests were analysed separately according to clinical breakpoints (CBP) defined by EUCAST version 7.0 and epidemiological cut off values (ECV). Etest as well as reference methods excellently recognised the WT isolates, which were susceptible to all tested triazoles, regardless of the method and CBP or ECV criteria used. The Etest recognized three non-WT isolates as resistant or intermediately sensitive to ITZ and POS and one as resistant to VOR. The categorical concordance between Etests and EUCAST and Etests and the CLSI method ranged from 90 to 100%. The interpretation of the results obtained from routine A. fumigatus Etests requires great caution. The use of the confirmative examinations with reference AST methods as well as with molecular tests is recommended.

**Key words**: *Aspergillus fumigatus*, triazole resistance, susceptibility testing, Etest, cyp51A sequence

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Abbreviations: ITZ, itraconazole; VOR, voriconazole; POS, posaconazole; CBP, clinical breakpoints; ECV, epidemiological cut off val¬ue; MIC, minimal inhibitory concentration; AST, antifungal susceptibility testing; CLSI, Clinical and Laboratory Standards Institute; ICU, intensive care unit; EUCAST, The European Committee on Antimicrobial Susceptibility Testing; CFU, colony forming unit; S, susceptible; R, resistant; I, intermediately susceptible; WT, wild-type; ATCC, American Type Culture Collection; ITS, internal transcribed spacer; TR, tandem repeats; EORTC, European Organization for Research and Treatment of Cancer.

## INTRODUCTION

Among fungi, Aspergillus plays an increasingly important role as a human and animal pathogen responsible for infections as well as allergies. Due to the ubiquitous character of fungal spores and its ability to easily disperse in the air, the respiratory tract is the most frequent port of entry and subsequently place of the disease development. The spectrum of antifungals suitable for treatment of aspergillosis includes triazole derivatives voriconazole (VOR), itraconazole (ITZ) and posaconazole (POS) as well as amphotericine B, representing polyenes. Aspergillus fumigatus, which currently is the most frequent filamentous fungus isolated from infection worldwide, shows an innate susceptibility to both triazoles and amphotericine B (Drgona et al., 2014). Taking into account pharmacokinetic properties and serious adverse effects connected with the implementation of amphotericine B, triazoles are recommended as a first line therapy for aspergillosis (Drgona et al., 2014; Groll et al., 2014; Maertens et al., 2011). An alarming phenomenon of acquired resistance of Aspergillus fumigatus to triazoles has recently been reported worldwide (Arendrup, 2014). Main mechanism of triazole resistance relies on mutations in target cyp 51A gene and/or in its promoter region and subsequently on substitution of amino acids in lanosterol 14 alpha-demethylase (protein CYP 51A). In Europe, 3.2% of clinical Aspergillus fumigatus isolates are resistant to triazoles and approximately 50% of them display mechanisms TR34/L98H (34bp tandem repeats incorporated in the promotor region and single point mutation of cyp 51A gene resulting in the substitution of leucine for histidine) (van der Linden et al., 2015). This mechanism is also frequent among environmental isolates (0-26%), which can be a source of infection in humans (Rodriguez-Tudela et al. 2008). This situation stresses the necessity to perform routine susceptibility tests. At present, two reference microdilution methods of susceptibility testing for filamentous fungi are available (Clinical and Laboratory Standards Institute, 2008; EUCAST DE-FINITIVÉ DOCUMENT E Def 9.2 2014). Nevertheless, due to the complicated procedures, they are difficult to perform in clinical laboratories. Instead, the suitability of Etests in routine diagnostics has been indicated, but its applicability to moulds is still under discussion (Espinel-Ingroff & Rezusta, 2002; Colosi *et al.*, 2012).

The aim of this study was to evaluate the accuracy of Etests used in routine diagnostics to examine the susceptibility of clinical isolates of *Aspergillus fumigatus* to triazoles (VOR, ITZ and POS). As a comparison, the reference microdilution methods according to the European Committee for Antimicrobial Susceptibility Testing (EUCAST) (Arendrup *et al.*, 2014) and Clinical and Laboratory Standards Institute (CLSI) (2008) were used. The susceptibility results were combined with the previously described sequences of the *cyp 51A* gene encoding lanosterol 14 alphademethylase – the target for triazoles (Nawrot *et al.*, 2017).

# MATERIAL AND METHODS

Strains and study design. Twenty clinical isolates of Aspergillus fumigatus from the laboratory collection of the Department of Microbiology, Medical University of War-saw (Poland) were selected for the study. The isolates were obtained in years 2007-2015 from patients with proven or probable aspergillosis according to EOTRC/ MSG criteria (each strain from another individual) (De Pauw et al., 2008). The patients were hospitalised in different departments including haematology - 6 patients, pulmonology - 6, ICU - 4, nephrology - 1, cardiosurgery -1, endocrinology – 1, and internal medicine – 1. Most isolates originated from respiratory tract samples (BAL - 8, sputum -5, broncho-aspirate -3) and the rest were from the pericardial fluid - 1 isolate, bone tissue - 1, blood -1, and wound -1. The strains were preserved by being frozen at -86°C using the Microbank<sup>TM</sup> Preservation System (Pro-Lab Diagnostics<sup>TM</sup>) and before testing were revived by subculture on Sabouraud dextrose agar. The isolates were identified as Aspergillus fumigatus based on the morphology and sequencing of rDNA and beta-tubuline genes. Seventeen isolates represented wild -type cyp51A sequence and three were positive for the mutation TR34/ L98H (Table 1) (Nawrot et al., 2017).

Each strain was tested for the susceptibility to itraconazole (ITZ), voriconazole (VOR), and posaconazole (POS) by microdilution methods according to EUCAST (Arendrup *et al.*, 2014) and CLSI (2008) and the obtained results were compared with Etest. In cases where significant discordance was observed between the applied methods, the Etests were performed again.

Susceptibility testing. Minimal Inhibitory Concentration (MIC) was determined with broth microdilution methods, according to EUCAST E.DEF 9.2 (Arendrup *et al.*, 2014) and according to the M38-A2 Document of the Clinical and Laboratory Standards Institute (2008). The RPMI1640 medium, buffered with MOPS and supplemented with glucose (2% in EUCAST and 0.2% in CLSI), was used to prepare inoculums of final density ranging  $1-2.5 \times 10^5$  CFU/mL (EUCAST) or  $0.4-5 \times 10^4$  CFU/mL (CLSI). The 100 µl aliquots of prepared cultures were inoculated to the flat-bottom wells of 96-wells microtiter plates containing 100 µl of tested drug solutions (two-fold serial drug dilutions ranging 0.0156–8 mg/L). The MIC was visually determined after 48 h of incubation at 35°C as the lowest drug concentration resulting in complete inhibition of fungal growth. All antifungals as well as main reagents (DMSO, RPMI 1640 medium, MOPS buffer) were obtained from Sigma-Aldrich.

The gradient susceptibility tests were performed using Etest-strips (BioMerieux, France) and RPMI 1640-2% glucose agar plates (Biomed, Poland) according to the manufacturer's instructions. Tests were incubated at 35°C for up to 48 h. The MIC values were indicated at the intersection of the inhibition zone with the Etest strips.

For the quality control of susceptibility tests, *Candida krusei* ATCC 6258 and *C. parapsilosis* ATCC 22019 were used.

The results of microdilution and Etests were analysed according to clinical breakpoints (CBP) defined by EU-CAST version 7.0 (http://www.eucast.org/fileadmin/ src/media/PDFs/EUCAST\_files/AFST/Antifungal\_ breakpoints\_v\_7.0.pdf ) and epidemiological cut off values (ECV) served as breakpoints before ratified clinical breakpoints by CLSI (Espinel-Ingroff et al., 2010). According to EUCAST the isolates of Aspergillus fumigatus were reported as resistant when the MIC value was higher than 2 mg/L for ITZ and VOR and 0.25 mg/L for POS, and as susceptible when the MIC value was equal or lower than 1 mg/L, 1 mg/L and 0.12 mg/L, respectively. Based on ECVs, the results of MIC classified the strains as resistant (non-WT) to ITZ, VOR, and POS when they exceeded 1 mg/L, 1 mg/L and 0.25 mg/L, respectively. The strains with MIC lower than or equal to 1 mg/L, 1 mg/L and 0.25 mg/L, respectively, were regarded as susceptible (WT).

## RESULTS

The results of susceptibility testing obtained by CLSI, EUCAST and Etest methods, as well as interpretation of susceptibility data and comparison of the methods are shown in Tables 1–3. The MIC values determined by CLSI and EUCAST methods did not differ in more than one dilution for VOR and POS and two dilutions for ITZ (Table 3). For most of the tested isolates, differences be-

Table 1. Results of susceptibility tests for Aspergillus fumigatus isolates with wild type and the mutation TR34/L98H in the cyp51A sequence

|  |                    |                   |                   | Ra                  | ange of MIC         | mg/L (mean)           |                       |                       |                       |
|--|--------------------|-------------------|-------------------|---------------------|---------------------|-----------------------|-----------------------|-----------------------|-----------------------|
| <i>cyp</i> 51A sequence (num-<br>ber of strains) | Itraconazole       |                   |                   |                     | Voriconazol         | e                     | Posaconazole          |                       |                       |
|  | CLSI               | EUCAST            | Etest             | CLSI                | EUCAST              | Etest                 | CLSI                  | EUCAST                | Etest                 |
| WT (n=17)  | 0.06–0.5<br>(0.26) | 0.061-1<br>(0.48) | 0.006-1<br>(0.28) | 0.125–0.5<br>(0.33) | 0.125–0.5<br>(0.28) | 0.016–0.38<br>(0.113) | 0.015–0.06<br>(0.058) | 0.015–0.06<br>(0.036) | 0.008–0.125<br>(0.04) |
| TR34/L98H positive<br>(n=3):                     |                    |                   |                   |                     |                     |                       |                       |                       |                       |
| <sup>1</sup> No 5287                             | >8                 | >8                | 1.5               | 2.0                 | 1.0                 | 1                     | 0.5                   | 0.5                   | 0.38                  |
| No 2708  | >8                 | >8                | 16                | 4.0                 | 2.0                 | 2.0                   | 0.5                   | 0.5                   | 0.5                   |
| <sup>1</sup> No 1473                             | >8                 | >8                | 3                 | 1.0                 | 2.0                 | 0.75                  | 0.25                  | 0.5                   | 0.38                  |

<sup>1</sup>results of Etests after 48 h incubation are presented; MICs obtained after 24 h incubation for ITR, VOR, POS were 0.5, 0.125,0.125 mg/L, respectively

| Table 2. Categorisation | of susceptibility | results based | on epidemiological | cut-off values | (ECV) and | clinical breakpoints | (CBP) of EU- |
|-------------------------|-------------------|---------------|--------------------|----------------|-----------|----------------------|--------------|
| CAST v7                 |                   |               |                    |                |           |                      |              |

|                                     | Categorisation according to ECV/CBP |        |       |              |        |       |              |        |       |
|-------------------------------------|-------------------------------------|--------|-------|--------------|--------|-------|--------------|--------|-------|
| cyp51A sequence (number of strains) | ltraconazole                        |        |       | Voriconazole |        |       | Posaconazole |        |       |
| -                                   | CLSI                                | EUCAST | Etest | CLSI         | EUCAST | Etest | CLSI         | EUCAST | Etest |
| WT (n=17)                           | S/S                                 | S/S    | S/S   | S/S          | S/S    | S/S   | S/S          | S/S    | S/S   |
| TR34/L98H positive (n=3):           |                                     |        |       |              |        |       |              |        |       |
| <sup>1</sup> No 5287                | R/R                                 | R/R    | R/I   | R/I          | S/S    | S/S   | R/R          | R/R    | R/R   |
| No 2708                             | R/R                                 | R/R    | R/R   | R/R          | R/I    | R/I   | R/R          | R/R    | R/R   |
| <sup>1</sup> No 1473                | R/R                                 | R/R    | R/R   | S/S          | R/I    | S/S   | S/I          | R/R    | R/R   |

<sup>1</sup>interpretation of Etests according to results obtained after 48 h; categorisation according to MICs obtained after 24 h was: S/S for ITZ, VOR, as well as POS

tween MICs obtained with Etests and CLSI or EUCAST were not higher than two dilutions (except of five, six and one isolate, which showed a difference of >2 dilutions for ITZ, VOR, or POS, respectively). Most isolates (17/20)showed wild-type cyp51A sequence and displayed susceptibility to all tested triazoles, regardless of the method used and of CLSI or EUCAST interpretative criteria. Some discrepancies in susceptibility results were found for three isolates, which were found to be positive for the mutation TR34/L98H. The MIC values obtained in reference and gradient tests for ITZ and POS against non-WT isolates were more similar than those obtained for VOR. Namely, the MIC of ITZ measured with CLSI and EUCAST was >8 mg/L for isolates No 5287, 1473 and 2708, whereas the results of the Etest were 1.5, 3, and 16 mg/L, respectively. Applying the CBP (EUCAST) allowed identification of one small categorical error (I instead of R) between results of the Etest and both reference methods. The MIC of POS for the isolate No1473 obtained with the CLSI method (0.25 mg/L) was lower than with EUCAST and the E-test (0.5 mg/L), and as a consequence the categorical disagreement of the major (S vs R, according to ECV) or minor (S vs I, according to CBP) level was found. In the case of VOR, the concordant result of CLSI and EUCAST was found for one non-WT isolate (No 2708), whereas for isolates No 1473 and No 5287, the MIC value was 1.0 mg/L in the CLSI and EUCAST method, respectively (Table 1). In contrast, the Etest detected resistance of one strain (No 2708). However, when compared to results obtained by CLSI or EUCAST, the Etests were in agreement with results obtained for 2/3 isolates. The overall categorical agreement between CLSI and EUCAST was 100% and 95% for ITZ and POS and 85-90% for VOR (depending on ECV or CBPs criterion). The concordance between Etests and EUCAST and Etests and the CLSI method ranged from 90 to 100%. (Table 3).

# DISCUSSION

Due to the complex morphological structure, moulds represent a difficult subject for laboratory examination. However, the development of the reference methods of susceptibility testing for filamentous fungi by CLSI and EUCAST represents major progress in mycological diagnostics. Despite the methods appearing to be similar (they varied only in the glucose concentration in the RPMI1640 medium and density of fungal inoculum), obtained results are sometimes inconsistent and the interpretative criteria must be developed individually for each method. The present study showed that although the correlation between the CLSI and EUCAST methods was very good (difference in  $\leq 2$  dilutions for 100% isolates), the interpretation according to the CBPs published by EUCAST may differ substantially from those based on epidemiological cut-off values used by CLSI. The extensive comparison of the EUCAST and CLSI methods for testing activity of VOR, POS, and ITZ against Aspergillus spp was described by Pfaller and coworkers (2011). Similarly to our study, the authors found excellent agreement between the methods and obtained MIC results did not differ in more than 2 dilutions for 100-97.5% of Aspergillus fumigatus isolates. Pfaller and coworkers (2011) categorised isolates as WT and non-WT, and found the highest number of discrepant results for POS. In the present study, applying the above mentioned criterion allowed us to find two major discrepancies (S vs. R) for VOR and one for POS.

Currently, a very important problem for routine diagnostics is the reliability and categorisation of the results obtained by commercial tests, especially the gradient diffusion Etests. Because the Etest does not require advanced equipment and is easy to perform it represents very attractive alternative for the microdilution method in many diagnostic laboratories. In this study, we evaluated the results of the

Table 3. Concordance in susceptibility results obtained with Etest and microdilution methods

|     | Number of strains with MIC differing in ±0/1/2/>2 dilution* |                         | Categorical agreement according to ECVs |                          |                         | Categorical agreement according to CBP |                   |                         |                           |
|-----|---|-------------------------|---|--------------------------|-------------------------|--|-------------------|-------------------------|---------------------------|
|     | CLSI <i>vs</i><br>EUCAST                                    | Etest <i>vs</i><br>CLSI | Etest <i>vs</i><br>EUCAST               | CLSI <i>vs</i><br>EUCAST | Etest <i>vs</i><br>CLSI | Etest vs<br>EUCAST                     | CLSI vs<br>EUCAST | Etest <i>vs</i><br>CLSI | Etest <i>vs</i><br>EUCAST |
| ITZ | 7/10/3/0  | 9/4/2/5                 | 4/6/5/5                                 | 100%                     | 100%                    | 100%                                   | 100%              | 95%                     | 95%                       |
| VOR | 12/8/0/0  | 3/9/2/6                 | 4/7/3/6                                 | 18/20 (90%)              | 18/20<br>(90%)          | 19/20<br>(95%)                         | 17/20<br>(85%)    | 18/20<br>(90%)          | 18/20<br>(90%)            |
| POS | 7/13/0/0  | 9/5/5/1                 | 9/8/2/1                                 | 19/20 (95%)              | 19/20<br>(95%)          | 100%                                   | 19/20<br>(95%)    | 19/20<br>(95%)          | 100%                      |

\*the values of MIC by Etests were adjusted to two-fold dilutions used in the reference methods

phenotypical antifungal susceptibility tests on a population of isolates with the defined sequence of the app51A gene. The isolates with the wild-type cyp51A sequence are likely to be susceptible to triazoles. The mutation TR34/L98H, which was detected in three isolates, represents the most frequent mechanism of azole-resistance in A. fumigatus and it is thought to be connected with cross-resistance to the Apergillus- active triazoles (Arendrup, 2014; Rodriguez-Tudela et al., 2008; van der Linden et al., 2015).

We found that ITZ and POS better indicate the TR34/ L98H - positive isolates than VOR. Some isolates harbouring this mutation retain "in vitro" susceptibility to VOR, which in clinical practices may lead to overlooking the resistance. These data support the general rule described by Rodriguez-Tudela et al. (2008), that currently known A. fumigatus susceptible to ITR do not show resistance to other triazoles, and that routine antifungal susceptibility tests should always include ITZ. On the other hand, the resistance to ITZ could be a result of other mechanisms, e.g., upregulation of efflux pumps and does not necessarily in-dicate cross-resistance (Rodriguez-Tudela et al., 2008; Shapiro et al., 2011; Brillowska-Dabrowska et al., 2015). Another important observation described in the presented study is that the MIC values obtained in the Etest were often lower than in the reference methods and that an incubation time of 48hrs for the Etest improved detection of resistance. An impact of incubation time on the tests results was also discussed by other authors. According to Espinel-Ingroff & Rezusta (2002) the 48 h incubation time of posaconazole Etests resulted in a better concordance with the reference method (95%), while a better agreement for itraconazole MICs was after 24 h (90.3%). Due to a low number of isolates included in this study we were not able to reliably validate compared methods, however, observed disagreements indicate the necessity of farther studies in this field. The general agreement between Etests and the reference method is high. Nevertheless, taking into account some inconsistent results obtained for resistant isolates, the confirmative examinations with reference AST methods as well as with molecular tests are recommended.

# **Conflict of interest**

U. Nawrot: received lecture honoraria and financial support for the participation in the congresses from Pfizer Polska Sp. z o.o.

All other authors declare that they have no conflict of interest.

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