



Article Fusarium Keratitis in Taiwan: Molecular Identification, Antifungal Susceptibilities, and Clinical Features

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Abstract: We performed molecular identification and antifungal susceptibilities of pathogens and investigated clinical features of 43 culture-proven *Fusarium* keratitis cases from 2015–2020 in Taiwan. The pathogens were identified by sequencing of their internal transcribed spacer regions of ribosomal DNA and translation elongation factor 1α gene; their antifungal susceptibilities (to seven agents) were determined by broth microdilution method. We also collected clinical data to compare the drug susceptibilities and clinical features of *Fusarium solani* species complex (FSSC) isolates with those of other *Fusarium* species complexes (non-FSSC). The FSSC accounted for 76.7% pathogens, among which *F. falciforme* (32.6%) and *F. keratoplasticum* (27.9%) were the most common species. Among clinically used antifungal agents, amphotericin B registered the lowest minimal inhibitory concentration (MIC), and the new azoles efinaconazole, lanoconazole and luliconazole, demonstrated even lower MICs against *Fusarium* species. The MICs of natamycin, voriconazole, chlorhexidine, lanoconazole, and luliconazole were higher for the FSSC than the non-FSSC, but no significant differences were noted in clinical outcomes, including corneal perforation and final visual acuity. In Taiwan, the FSSC was the most common complex in *Fusarium* keratitis; its MICs for five tested antifungal agents were higher than those of non-FSSC, but the clinical outcomes did not differ significantly.

Keywords: Fusarium keratitis; molecular identification; antifungal susceptibility

1. Introduction

Fusarium species are the most frequent causative agent of fungal keratitis, a corneal infection that can lead to severe vision loss [1,2]. *Fusarium* keratitis risk factors include contact lens use, ocular trauma, ocular surgery, topical steroid use, and immunosuppression [3]. With increasing numbers of contact lens wearers, *Fusarium* keratitis is becoming a critical health concern worldwide. In 2005, an outbreak of contact lens-associated *Fusarium* keratitis was reported in multiple regions, including Hong Kong, Singapore, Europe, and the United States (US) [4–6]. This event was reportedly associated with decreased antimicrobial activity in Bausch and Lomb's contact lens cleaning solution ReNu with MoistureLoc [7]. *Fusarium* keratitis is an important issue in multiple countries, and Taiwan is no exception. *Fusarium* species account for 44.6% of the filamentous fungal keratitis in Taiwan [3], and the proportion of *Fusarium*-related cases among all microbial keratitis increased from 4.0% to 6.4% between 1992 and 2016 [8].



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Copyright: © 2022 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (https:// creativecommons.org/licenses/by/ 4.0/). Early diagnosis and prompt delivery of antifungal agents are key factors in *Fusarium* keratitis treatment. Although traditional morphology is frequently adopted in daily clinical practice for diagnosis, it is time-consuming and cannot be used to differentiate isolates to the species level. In contrast, molecular diagnosis has advantages of expediency, accuracy, and genotyping capacity. Molecular identification is crucial to epidemiology and can influence therapy and outcomes in cases of interspecies differences in antifungal susceptibilities or virulence. However, despite those advantages, molecular diagnosis was still not widely used around the world. From a recent meta-review, only 11.9% of the *Fusarium* isolates (n = 628) were identified to species level among patients with *Fusarium* keratitis (n = 5294) [9].

Epidemiological studies involving molecular diagnosis of *Fusarium* keratitis have been conducted in several countries, including Germany, the Netherlands, India, and Brazil [10–13]. However, no such studies have been performed in East Asia. Because the epidemiological patterns may vary by country and area, we performed molecular identification and antifungal susceptibilities of *Fusarium* species in Taiwan. In addition, studies of the correlations of *Fusarium* species and antifungal susceptibilities with clinical outcomes are rare; therefore, we investigated such correlations by comparing members of the *F. solani* species complex (FSSC) with those of other *Fusarium* (i.e., non-FSSC) species.

2. Materials and Method

2.1. Study Population and Data Collection

We extracted information regarding culture-proven *Fusarium* keratitis cases from 1 January 2015, through 31 December 2020, from the database of the microbiological laboratory at CGMH, Linkou branch, Taiwan. We obtained cornea scrapings from patients with infectious keratitis and sent specimens to smear and culture examination. The cornea scraping specimens were cultivated on blood and chocolate agar, modified Sabouraud agar, Lowenstein–Jensen agar slants, and thioglycolate broth. Positive *Fusarium* culture was defined as *Fusarium* species growth identified through morphology on two media, fungal elements observed in smears and *Fusarium* species growth on one medium, or confluent growth of *Fusarium* species on one medium.

We reviewed the medical charts of the patients associated with the keratitis cases for demographic data, systemic and ocular disease history, predisposing factors, initial ocular presentation, antifungal treatment, surgery requirement, hospitalization, corneal perforation or endophthalmitis, and initial and final visual acuity (VA). The predisposing factors were ocular trauma, an outdoor occupation or gardening habit, contact lens use, topical steroid use, preexisting ocular disease, and recent ocular surgery. Preexisting ocular disease referred to any disease that interfered with the epithelial integrity of the cornea. Recent ocular surgery meant any surgery performed up to 3 months before the ulcer. An ulcer was defined as central if it was less than 2 mm from the corneal center and peripheral if it extended to within 2 mm of the limbus; otherwise, the ulcer was defined as paracentral if its location was between the definition of cornea center and periphery. If the epithelium defect covered the cornea from center to periphery in all four quadrants, it was defined as near total epithelium defect. Ulcers less than 2 mm in diameter were considered small, those 2–6 mm in diameter were considered medium, and those larger than 6 mm were considered large. As a clinical outcome, corneal perforation also included cases of impending perforation that required therapeutic penetrating keratoplasty (TPK) or amniotic membrane transplantation (AMT). VA was measured using Snellen charts and converted to the logarithm of minimum angle of resolution (logMAR) VA. As suggested by Schulze-Bonsel et al. [14], nonnumerical VA data were converted to logMAR values as follows: counting fingers = 2, hand motion = 2.3, light perception = 2.7, and no light perception = 3.0. Initial VA was measured on the first visit for *Fusarium* keratitis. Final VA, if available, was measured as best corrected VA (BCVA) on the final visit.

2.2. DNA Extraction, Amplification, and Sequencing

All the *Fusarium* isolates underwent molecular diagnosis process. If one patient received multiple times of corneal fungal culture during the treatment period, the *Fusarium* isolates each time would all undergo molecular diagnosis individually and be given their own strain number.

Fungal isolates were subcultured on potato dextrose agar for purification. Thereafter, part of the mycelium was collected and placed in a plastic vial, to which 0.13 g of metal beads and 800 µL of lysis buffer (Tris buffer, surfactants, pH 8.0) were added. The vial was then transferred to a cell disruptor (Mini-BeadBeater 16, BioSpec, Bartlesville, OK, USA) to break down the cell walls. The buffer fluid containing the fungal fragments was transferred to a DNA extraction kit, and genomic DNA was extracted with a Smart LabAssist (TANBead, Taiwan) automatic DNA extraction system. Internal transcribed spacers of ribosomal DNA (ITS) and translation elongation factor-1 α gene (*TEF*-1 α) were used for molecular identification. The ITS regions were amplified with primers ITS1 (TCCGTAG-GTGAACCTGCGG) and ITS4 (TCCTCCGCTTATTGATATGC), and TEF-1 α gene was amplified with primers EF1 (ATGGGTAAGGARGACAAGAC) and EF2 (GGARGTACCAGT-SATCATG). The PCR conditions were as described previously [15,16]. The PCR products were confirmed by gel electrophoresis, then purified and submitted for DNA sequencing with ABI Prism 3730 xl DNA analyzer (Applied Biosystems, Foster City, CA, USA). The sequences were uploaded to Fusarium MLST database at Mycobank website (https://fusarium.mycobank.org/, accessed on 1 February 2022) and Fusarium Database (http://isolate.fusariumdb.org, accessed on 1 February 2022) for preliminary species identification. All sequences generated in this study were uploaded to the DNA Data Bank of Japan (DDBJ) [17].

2.3. Phylogenetic Analyses

Based on the preliminary identification results from the *Fusarium* MLST database, the ITS and *TEF-1* α sequences of allied *Fusarium* species were retrieved from GenBank (Table 1). Each region was first aligned with MAFFT online version [18], and poorly aligned regions were removed manually or with Gblocks online software [19]. DNA evolution models were determined with Smart Model Selection (SMS) [20]. The two loci were then concatenated for the following phylogenetic analyses. Maximum likelihood trees were inferred by RAxML-NG v. 1.1.0, and Felsenstein's bootstrap (FBP) statistical support was calculated from 1000 resample data sets [21]. Bayesian inference tree was inferred by using MrBayes v3.2.6 x64 [22]. The analysis was performed with two MCMC chains for 1,000,000 generations, and one tree was sampled for every 1000 generations. After discarding the first 250 trees (burn-in), a consensus tree was obtained from the remaining 750 trees. All analyses were conducted on the Linux Mint 20.3 (64-bit) operating system, and to ensure reproducibility, the random seeds were explicitly set to 56 wherever they were needed.

2.4. Antifungal Susceptibility Testing

Broth microdilution was used to determine the minimal inhibitory concentration (MIC) of antifungals to various strains. The procedures were performed in accordance with the third edition of *M38: Reference Method for Broth Dilution Antifungal Susceptibility Testing of Filamentous Fungi*, published by the Clinical and Laboratory Standards Institute [23]. Pure powders of antifungals and chemical for testing were purchased from Sigma-Aldrich[®] and the range of concentration was amphotericin B (16–0.031 µg/mL), natamycin (32–0.063 µg/mL), voriconazole (16–0.031 µg/mL), chlorhexidine (256–0.5 µg/mL), efinaconazole (4–0.008 µg/mL), lanoconazole (0.5–0.001 µg/mL), and luliconazole (0.5–0.001 µg/mL). ATCC 22,019 *Candida parapsilosis* and ATCC 6258 *Candida krusei* were used as control. MIC endpoints were determined using a reading mirror after 48 h of incubation at 35 °C and indicated by a 100% inhibition of growth compared with drug-free growth control well for all the drugs and chemicals.

Fusarium Isolates	Species	Accession Number					
Strain Number	Species	ITS	TEF-1α				
CGMHD0184	Fusarium keratoplasticum	LC683270	LC683322				
CGMHD0412	Fusarium falciforme	LC683271	LC683323				
CGMHD0604	Fusarium falciforme	LC683272	LC683324				
CGMHD0633	Fusarium suttonianum	LC683273	LC683325				
CGMHD0634	Fusarium falciforme	LC683274	LC683326				
CGMHD0820	Fusarium keratoplasticum	LC683275	LC683327				
CGMHD0863	Fusarium breve	LC683276	LC683328				
CGMHD0998	Fusarium keratoplasticum	LC683277	LC683329				
CGMHD1000	Fusarium keratoplasticum	LC683278	LC683330				
CGMHD1102	Fusarium falciforme	LC683279	LC683331				
CGMHD1463	Fusarium falciforme	LC683280	LC683332				
CGMHD1481	Fusarium falciforme	LC683281	LC683333				
CGMHD1606	Fusarium falciforme	LC683282	LC683334				
CGMHD1794	Fusarium keratoplasticum	LC683283	LC683335				
CGMHD1795	Fusarium falciforme	LC683284	LC683336				
CGMHD1816	Fusarium falciforme	LC683285	LC683337				
CGMHD1817	Fusarium falciforme	LC683286	LC683338				
CGMHD1982	Fusarium keratoplasticum	LC683287	LC683339				
CGMHD2012	Fusarium silvicola	LC683288	LC683340				
CGMHD2013	Fusarium falciforme	LC683289	LC683341				
CGMHD2045	Fusarium keratoplasticum	LC683290	LC683342				
CGMHD2046	Fusarium keratoplasticum	LC683291	LC683343				
CGMHD2064	Fusarium keratoplasticum	LC683292	LC683344				
CGMHD2065	Fusarium keratoplasticum	LC683293	LC683345				
CGMHD2066	Fusarium keratoplasticum	LC683294	LC683346				
CGMHD2067	Fusarium keratoplasticum	LC683295	LC683347				
CGMHD2068	Fusarium keratoplasticum	LC683296	LC683348				
CGMHD2086	Fusarium keratoplasticum	LC683297	LC683349				
CGMHD2087	Fusarium keratoplasticum	LC683298	LC683350				
CGMHD2134	Fusarium keratoplasticum	LC683299	LC683351				
CGMHD2214	Fusarium keratoplasticum	LC683300	LC683352				
CGMHD2224	Fusarium suttonianum	LC683301	LC683353				
CGMHD2402	Fusarium falciforme	LC683302	LC683354				
CGMHD2521	Fusarium falciforme	LC683303	LC683355				
CGMHD2740	Fusarium suttonianum	LC683304	LC683356				
CGMHD2994	Fusarium falciforme	LC683305	LC683357				
CGMHD3072	Fusarium keratoplasticum	LC683306	LC683358				
CGMHD3140	Fusarium keratoplasticum	LC683307	LC683359				
CGMHD3224	Fusarium keratoplasticum	LC683308	LC683360				
CGMHD3255	Fusarium falciforme	LC683309	LC683361				
CGMHD3337	Fusarium falciforme	LC683310	LC683362				
CGMHD4138	Fusarium suttonianum	LC683311	LC683363				
CGMHD0275	Fusarium veterinarium	LC683312	LC683364				
CGMHD1515	Fusarium nirenbergiae	LC683313	LC683365				
CGMHD3336	Fusarium lacertarum	LC683314	LC683366				
CGMHD2044	Fusarium sp. FIESC ^a	LC683315	LC683367				
CGMHD1650	Fusarium aff. circinatum	LC683316	LC683368				
CGMHD3368	Fusarium annulatum	LC683317	LC683369				
CGMHD1660	Fusarium sp. FDSC ^b	LC683318	LC683370				
CGMHD2001	Fusarium delphinoides	LC683319	LC683371				
CGMHD2436	Fusarium sp. FDSC ^b	LC683320	LC683372				
CGMHD3049	<i>Fusarium</i> sp. FDSC ^b	LC683321	LC683373				
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Table 1. *Fusarium* species and GenBank accession numbers of *Fusarium* isolates ($n = 52^*$) from patients with keratitis.

* Four out of the 43 patients received multiple corneal fungal culture during the treatment period, so eventually, 52 *Fusarium* isolates were collected. All the 52 isolates underwent molecular diagnosis and antifungal susceptibility tests individually. ITS: internal transcribed spacer region; *TEF-1* α : translation elongation factor -1 α . ^a The *TEF-1* α

similarity of this isolate was only 96.3% to an isolate of *F. incarnatum* in Genbank (MAFF 236386), but in another article, the same species was renamed as *F. semitectum* [25]. Hence, further species identification was required for this isolate. ^b Three isolates (CGMHD 1660, 2436, and 3049) were clustered with NRRL 37,393 and CBS 110,307 in *F. dimerum* species complex (FDSC), and these strains had already been identified by Hans-Josef Schroers et al., in 2009 but it still remained unnamed [24].

If one patient received corneal fungal culture multiple times and two or more isolates were obtained, each *Fusarium* isolate was subjected to antifungal susceptibility tests individually. However, we only used the drug susceptibility results from the first *Fusarium* isolates when presenting the antifungal susceptibilities or comparing the MIC between different *Fusarium* species complex.

2.5. Statistical Analysis

The isolates were separated by molecular identification into the FSSC and non-FSSC groups. The antifungal susceptibilities and clinical features of these two groups were compared. The initial and final logMAR VA values were compared using Mann–Whitney U tests. Categorical demographic and outcome variables were compared using chi-square tests; continuous variables in demographic data, clinical outcomes, and MIC results were compared using Student's t test. Statistical significance was defined as p < 0.05. All statistical analyses were performed using SPSS version 25 (IBM, New York, NY, USA).

3. Results

3.1. Molecular Identification of Fusarium Isolates

A total of 43 cases of culture-proven *Fusarium* keratitis from the study period of 2015–2020 were analyzed. Four out of the 43 patients received multiple corneal scrapings for fungal culture during the treatment period, so eventually, 52 *Fusarium* isolates were obtained and underwent molecular diagnosis. Based on multilocus phylogenetic analyses, 14 *Fusarium* species, belonging to five species complexes, were identified and the result is shown in Table 1 and Figure 1. For those four patients receiving multiple fungal culture, their pathogens remained the same throughout the sampling period.

Three isolates (CGMHD 1660, 2436, and 3049) were clustered with NRRL 37,393 and CBS 110,307 in *F. dimerum* species complex (FDSC), and these strains had already been identified by Hans-Josef Schroers et al., in 2009, but it still remained unnamed. Hence, we presented these three isolates as *Fusarium* sp. FDSC in this article [24]. The species of CGMHD 2044 isolate was still uncertain, because its *TEF-1* α result had only 96.3% similarity to an isolate of *F. incarnatum* in Genbank (MAFF 236386), but in another article, the same species was renamed as *F. semitectum* [25]. Hence, further species identification was required for this isolate and it was only presented as *Fusarium* sp. FIESC in this article.

The pathogen distribution in 43 patients with keratitis based on molecular identification is shown in Table 2. Thirty-three cases (76.7%) belonged to the FSSC group, among which 14 (32.6%) were *F. falciforme*, and 12 (27.9%) were *F. keratoplasticum*. Four cases (9.3%) belonged to the FDSC, among which three (7.0%) were *F. dimerum* SC and one (2.3%) was *F. delphinoides*. The *F. oxysporum* species complex (FOSC), *F. incarnatum-equiseti* species complex (FIESC), and *F. fujikuroi* species complex (FFSC) each accounted for two cases (4.7%). For data analysis, those cases were divided into the FSSC and non-FSSC groups.

F. lacertarum NRRL 36123



Figure 1. The phylogenetic tree of the 52 Fusarium isolates from patients with keratitis. All the Fusarium isolates were labelled in bold and strain number were provided.

Species Complex Number (%)	Fusarium Species	Case Number (%)					
	F. falciforme	14 (32.6%)					
	F. keratoplasticum	12 (27.9%)					
FSSC	F. solani.	3 (7.0%)					
33 (76.7%)	F. suttonianum	2 (4.7%)					
	F. silvicola	1 (2.3%)					
	F. breve	1 (2.3%)					
FDSC	Fusarium sp. FDSC ^a	3 (7.0%)					
4 (9.3%)	F. delphinoides	1 (2.3%)					
FOSC	F. nirenbergiae	1 (2.3%)					
2 (4.7%)	F. veterinarium	1 (2.3%)					
FIESC	F. lacertarum	1 (2.3%)					
2 (4.7%)	<i>Fusarium</i> sp. FIESC ^b	1 (2.3%)					
FFSC	F. aff. circinatum	1 (2.3%)					
2 (4.7%)	F. annulatum	1 (2.3%)					

Table 2. Fusarium species distribution in patients with keratitis.

FSSC: *Fusarium solani* species complex; FDSC: *Fusarium dimerum* species complex; FOSC: *Fusarium oxysporum* species complex; FIESC: *Fusarium incarnatum-equiseti* species complex; FFSC: *Fusarium fujikuroi* species complex. ^a Hans-Josef Schroers et al., had identified this *Fusarium* species in 2009 but it was still unnamed [24]. ^b Further species identification was required for this isolate because its *TEF-1a* similarity was only 96.3% to an isolate of *F. incarnatum* in Genbank (MAFF 236386), but in another article, the same species was renamed as *F. semitectum* [25].

3.2. Antifungal Susceptibilities

As indicated in Table 3, we compared the geometric mean (GM) MICs of each antifungal agent against the FSSC and non-FSSC isolates. Among the antifungal agents currently used in clinical practice (i.e., amphotericin B, natamycin, voriconazole, and chlorhexidine), amphotericin B had the lowest GM MIC against both the FSSC and non-FSSC isolates, but the new azoles, including efinaconazole, lanoconazole, and luliconazole, demonstrated even lower GM MICs for *Fusarium* species. The GM MICs of natamycin (p = 0.021), voriconazole (p = 0.010), chlorhexidine (p < 0.001), lanoconazole (p = 0.004), and luliconazole (p = 0.026) were higher for the FSSC isolates than the non-FSSC isolates. In addition, the FSSC group also exhibited a trend of higher MIC in efinaconazole (p = 0.052). The detailed MIC data for each isolate are presented in the Appendix A, Table A1.

3.3. Demographic Data, Predisposing Factors, and Initial Presentation

As listed in Table 4, the study sample comprised cases from 29 (67.4%) men and 14 (32.6%) women. Their average age was 51.5 ± 19.5 (11–91) years. No differences between the FSSC and non-FSSC groups were observed in terms of age or sex. Outdoor occupation or gardening habit (n = 23, 53.5%) and ocular trauma (n = 19, 44.2%) were the most common predisposing factors, followed by recent ocular surgery (n = 5, 11.6%), contact lens use (n = 4, 9.3%), preexisting ocular disease (n = 4, 9.3%), and topical steroid use (n = 1, 2.3%). The proportion of contact lens use was significantly higher in the non-FSSC group than in the FSSC group (30% vs. 3%, p = 0.010), but no significant differences were noted in the other predisposing factors. Most of the corneal ulcers were in the paracentral area (n = 27, 62.8%) and of medium size (n = 27, 62.8%). No significant differences in ulcer area or location were observed between the FSSC and non-FSSC groups. At initial presentation, 23 of the case patients (53.5\%) had hypopyon, but none exhibited corneal perforation.

Antifungal Agents	Fusarium Species	Range of MIC (mg/L)	GM MIC (mg/L)	p Value
Amphotericin B	FSSC Non-FSSC	0.5~4 1~2	1.66 1.36	0.263
Natamycin	FSSC Non-FSSC	4~16 2~8	5.79 3.70	0.021
Voriconazole	FSSC Non-FSSC	2~16 2~8	7.29 2.94	0.010
Chlorhexidine	FSSC Non-FSSC	16~32 8~16	18.38 9.33	<0.001
Efinaconazole	FSSC Non-FSSC	0.5~2 0.25~4	1.15 0.46	0.052
Lanoconazole	FSSC Non-FSSC	0.125~0.5 0.063~0.25	0.20 0.09	0.004
Luliconazole	FSSC Non-FSSC	0.031~0.25 0.016~0.063	0.06 0.03	0.026

Table 3. Comparison of antifungal susceptibilities between FSSC (n = 33) and non-FSSC (n = 10).

FSSC: Fusarium solani species complex; GM: geometric mean; MIC: minimal inhibitory concentration; non-FSSC: non-Fusarium solani species complex.

Table 4. Demographic data, predisposing factors, and initial presentation of 43 patients with *Fusarium* keratitis.

		Total	(%)	FSSC	2 (%)	Non-FS	<i>p</i> -Value	
	Total case number	43		33		10		
Gender	Male	29	(67.4%)	23	(69.7%)	6	(60.0%)	0.566
	Female	14	(32.6%)	10	(30.3%)	4	(40.0%)	
Age	Average (range)	51.5 ± 19.5	(11~91)	50.5 ± 20.0	(11~91)	54.6 ± 18.4	(19~86)	0.572
	Outdoor occupation or gardening habit	23	(53.5%)	18	(54.5%)	5	(50.0%)	0.801
	Trauma	19	(44.2%)	17	(51.5%)	2	(20.0%)	0.079
Predisposing	Recent ocular surgery	5	(11.6%)	4	(12.1%)	1	(10.0%)	0.855
Factors ^a	Contact lens use	4	(9.3%)	1	(3.0%)	3	(30.0%)	0.010
	Preexisting ocular disease	4	(9.3%)	3	(9.1%)	1	(10.0%)	0.931
	Topical steroid use	1	(2.3%)	1	(3.0%)	0	(0.0%)	0.578
	Central	5	(11.6%)	4	(12.1%)	1	(10.0%)	0.855
	Paracentral	27	(62.8%)	21	(63.6%)	6	(60.0%)	0.835
Ulcer location	Peripheral	7	(16.3%)	4	(12.1%)	3	(30.0%)	0.180
	Near total	4	(9.3%)	4	(12.1%)	0	(0.0%)	0.248
	Small	7	(16.3%)	5	(15.2%)	2	(20.0%)	0.716
Ulcer area	Medium	27	(62.8%)	20	(60.6%)	7	(70.0%)	0.59
	Large	9	(20.9%)	8	(24.2%)	1	(10.0%)	0.332
Hypopyon		23	(53.5%)	19	(57.6%)	4	(40.0%)	0.878

^a Total percentage is more than 100% because 18 patients had multiple risk factors. FSSC: *Fusarium solani* species complex; non-FSSC: *non- Fusarium solani* species complex.

3.4. Treatment and Outcomes

As listed in Table 5, among the 43 patients, 29 (66.7%) required hospitalization, for 12.3 \pm 15.3 days on average. All the patients were treated with antifungal agents, for an average 31.8 \pm 33.4 days. The most used topical antifungal agents were natamycin (44.2%), voriconazole (44.2%), and amphotericin B (39.5%), which were used either as single agents or in combination. Twenty-three patients (53.5%) underwent surgical procedures, including keratectomy (n = 12, 27.9%), AMT (n = 4, 9.3%), and TPK (n = 7, 16.3%). Two patients developed endophthalmitis (4.7%), and ten experienced corneal perforation (23.3%). Final VA was worse than 20/200 in 15 patients (34.9%). No significant differ-

Total (%) **FSSC (%)** Non-FSSC (%) p-Value 29 (66.7%) 22 (66.7%) 7 (67.4%) Hospitalization 13.4 ± 16.9 8.7 ± 7.0 average days 12.3 ± 15.3 0.402 Topical natamycin 19 (44.2%) 15 (45.5%) 4 (40.0%) 0.761 Topical voriconazole 19 (44.2%) 15 (45.5%)4 (40.0%)0.761 (42.4%) (39.5%) (30.0%) Topical amphotericin B 17 14 3 0.481 Medical (15.2%) 0 5 5 (0.0%) 0.075 (11.6%)Oral voriconazole Treatment^a Oral Fluconazole 2 (4.7%) 2 (6.1%) 0 (0.0%) 0.425 Antifungal usage 35.0 ± 36.7 21.8 ± 17.3 31.8 ± 33.4 0.282 (average days) (27.9%) 9 (27.3%) (30.0%) 0.866 Keratectomy 12 3 Surgical (9.3%) 3 (9.1%) (10.0%) 0.931 AMT 4 1 treatment 7 5 2 (20.0%)0.716 TPK (16.3%) (15.2%) Endophthalmitis 2 (4.7%) 1 1 0.359 (3.0%)(10.0%)Complications Perforation 10 (23.3%) 7 (21.2%) 3 (30.0%) 0.564Median initial logMAR 1.30 1.70 0.47 0.171 $(0.55 \sim 2.28)$ $(0.70 \sim 2.14)$ $(0.17 \sim 2.20)$ VA (IOR) Visual Median final logMAR outcomes $(0.28 \sim 1.75)$ 0.70 $(0.40 \sim 1.0)$ 0.40 $(0.22 \sim 1.61)$ 0.878 0.61 VA (IQR) Final VA < 20/200 15 (34.9%)12 (36.4%) 3 (30.0%)0.572

ences were observed between the FSSC and non-FSSC groups in treatment, complications, or visual outcomes.

Table 5. Treatment and outcomes of 43 patients with Fusarium keratitis.

^a Total percentage of treatment method exceeded 100% because some patients received multiple antifungal treatment or were combined with surgical treatment. AMT: amniotic membrane transplantation; IQR: interquartile range; TPK: therapeutic penetrating keratoplasty; VA: visual acuity.

4. Discussion

We investigated the epidemiological data of *Fusarium* isolates from 43 cases of *Fusarium* keratitis diagnosed at a referral center in Taiwan. We identified 13 *Fusarium* species belonging to five species complexes; 33 of the isolates (76.7%) belonged to the FSSC. We also tested susceptibilities to seven antifungals, among which amphotericin B achieved the lowest MICs. The GM MICs of natamycin, voriconazole, chlorhexidine, lanoconazole, and luliconazole were significantly higher for FSSC isolates than for non-FSSC isolates. An outdoor occupation or gardening habit and ocular trauma were the most common predisposing factors; 23 patients (53.5%) required surgical procedures, including ten (23.3%) operations for perforation, and 15 patients (34.9%) had final VA of less than 20/200. No significant differences in clinical features were observed between the FSSC and non-FSSC groups.

The FSSC accounted for more than three-quarters of the *Fusarium* keratitis cases, a finding consistent with those of epidemiological studies of *Fusarium* keratitis in other countries [10–13,26–28]. Although the FSSC was the leading cause of *Fusarium* keratitis, the proportion of FSSC-related keratitis was variable in different countries. It has been reported as high as 75–88% in tropical or subtropical areas such as India, Brazil, or Miami (US) [12,13,26]. By contrast, the proportion of FSSC-related keratitis cases reported in temperate countries such as Germany and the Netherlands has been 41–59% [10,11].

In our study, the most common predisposing factors were an outdoor occupation or gardening habit and ocular trauma; these accounted for approximately half of the cases. By contrast, contact lens use was a factor in only 9.3% of cases. Because our hospital is a regional referral center near industrial and agricultural areas, our results are similar to those reported in countries with large agricultural or industrial sectors, such as Mexico and Brazil, in which ocular trauma is the most common risk factor (57.4% and 48.8%, respectively) and contact lens is implicated in only 8.1% and 4.9% of cases, respectively [12,13]. By contrast, in high-income countries, such as the US, Germany, and the Netherlands, contact lens use is the primary risk factor and associated with 63.5–81.8% of cases, whereas ocular trauma is implicated in only 5.6–17% of cases [10,11,29].

So far, there is still no established clinical breakpoint for antifungal susceptibility test for Fusarium species according to European Committee on Antimicrobial Susceptibility Testing (EUCAST) [30]. In our study, amphotericin B registered the lowest GM MIC (FSSC = 1.66 mg/L, non-FSSC = 1.36 mg/L), followed by natamycin (FSSC = 5.79 mg/L), non-FSSC = 3.70 mg/L) and voriconazole (FSSC = 7.29 mg/L, non-FSSC = 2.94 mg/L) among the antifungal agents used to treat *Fusarium* keratitis. Many epidemiological studies have reported comparable results, with amphotericin B exhibiting the lowest mean MIC (1.0-2.0 mg/L); the mean MIC results for natamycin (4.0-6.5 mg/L) and voriconazole (5.0-9.9 mg/L) are also similar to our results [10,11,13,26,27,29]. In the US, Oechesler et al., compared the antifungal susceptibilities of FSSC and non-FSSC species and reported that the MIC of voriconazole was higher for FSSC than for non-FSSC species but observed no differences in the MICs for amphotericin B or natamycin [29], which contradicted with our findings that the MICs of natamycin and voriconazole were higher for the FSSC group but no differences between the MICs of amphotericin B against FSSC and non-FSSC species. The discrepancies in these results are probably due to regional differences or differences in the proportions of Fusarium species complexes (with FSSC and FOSC accounting for 75% and 16%, respectively, in the study of Oechesler et al.).

Fusarium keratitis-related information on molecular identification, antifungal susceptibilities, and correlations with clinical outcomes is scarce [13,26,27,29], but such information may be helpful for guiding treatment selection and predicting patient outcomes. According to an study in Miami (US), Oechesler et al. reported that FSSC cases were associated with longer recovery times, poorer follow-up BCVA, and a greater need for urgent surgical procedures than were non-FSSC cases; in addition, they suggested that other factors such as interspecies differences in pathogenicity, rather than antifungal susceptibilities, may explain the difference in outcomes between FSSC and non-FSSC case because no difference was observed between groups in the MICs of natamycin or amphotericin B—the most commonly used topical antifungal agents [26]. However, in another study in Brazil by Oechesler et al., they did not obtain the same findings except for the greater need for TPK in FSSC cases than in non-FSSC cases, which may have been associated with a higher natamycin MIC [13]. In our study, despite higher MICs of multiple antifungal agents for the FSSC group, such differences in clinical outcomes were not observed between groups. Again, regional differences or differences in the proportions of *Fusarium* species complexes may explain these discrepant results. In addition, our sample may have been too small for differences to become evident.

Although amphotericin B has had the lowest MIC against *Fusarium* species in multiple studies, no consensus has been reached regarding which antifungal agent is optimal for the treatment of *Fusarium* keratitis. Studies have suggested amphotericin B as the first choice treatment for yeast infection and aspergillosis and natamycin for filamentous fungal keratitis [1,31]. In addition, the absolute MIC can be converted in a relative MIC on the basis of the typically prescribed concentration, as suggested by Lalitha et al., revealing that amphotericin B has a higher relative MIC and may be less likely to be the optimal treatment for *Fusarium* keratitis [32]. For example, the typically prescribed concentration of amphotericin B is 1.5 mg/mL, which is 903 times the MIC for FSSC cases (1.66 mg/L). However, the concentration of natamycin in commercial eyedrops is 5%, which is 3012-fold the MIC for FSSC cases. Based on the typically prescribed concentration, the relative MIC of natamycin is markedly higher than that of amphotericin B. Although we cannot draw conclusions directly from relative MICs because drug bioavailability and corneal penetration must also be considered, the concept of relative MIC provides a new perspective on how antifungal agents operate in clinical practice. In a study of antifungal susceptibilities in fungal keratitis cases in Shandong Province, China, 94.2% of F. solani and 91.3% of F. oxysporum cases were susceptible to natamycin, whereas only 82.4% of F. solani and 74% of F. oxysporum cases were susceptible to amphotericin B [33]. Thus, the authors of that study suggested natamycin as the first choice for the treatment of Fusarium keratitis. Moreover, amphotericin B's ocular toxicity and strong irritation of the ocular surface limit its clinical

application. In our clinical experience, topical natamycin 5% is the optimal choice for treating Fusarium keratitis, but in cases where the clinical response is poor, we recommend adding amphotericin B or voriconazole. However, due to frequent natamycin shortages, we occasionally must begin treatment with amphotericin B or voriconazole instead, explaining why only 44.2% of cases were treated with natamycin (Table 5). The optimal first-line antifungal agent for filamentous fungal keratitis remains debated. As mentioned earlier, although amphotericin B is typically the first choice for conditions caused by Aspergillosis and yeast infections [1,31], its high relative MIC, ocular toxicity, and ocular surface irritation indicate that it may not be the optimal choice. Although voriconazole demonstrates broad-spectrum antifungal activity and favorable corneal penetration, the Mycotic Ulcer Treatment Trial revealed that patients with natamycin-treated fungal keratitis had better 3-month BCVA and a lower risk of corneal perforation than did those with voriconazoletreated cases. However, in Aspergillus keratitis, natamycin-treated cases are associated with greater risks of corneal perforation and TPK requirement than voriconazole-treated cases are [34]. Therefore, identifying a new first-line antifungal agent for filament fungal keratitis is essential. Chlorhexidine 0.2% is commonly used in Acanthamoeba keratitis treatment, but seldom used as an antifungal agent in clinical practice. A recent evidence-based study recommended that chlorhexidine 0.2% could be an alternative to natamycin in the treatment of filamentous fungal keratitis [35], when natamycin is not available. In Taiwan, we often encountered with shortage of natamycin, so we decided to investigate the chlorhexidine susceptibility in the *Fusarium* isolates. The GM MIC of chlorhexidine in our study was similar to that in the Dutch study by Oliveira dos Santos et al. [36]; they further demonstrated chlorhexidine having fungicidal activity against 90% of *F. oxysporum* strains and 100% of the F. solani strains, which supported the clinical efficacy of chlorhexidine. Moreover, we assessed susceptibilities to three new azoles, namely efinaconazole, lanoconazole, and luliconazole; all three drugs demonstrated extremely low MICs against all Fusarium species In a previous study, efinaconazole, lanoconazole, and luliconazole also registered low MICs against *Fusarium* isolates from corneal scrapings in vitro, with MIC50 values of 1, 0.06, and 0.03 mg/L, respectively [37]. In addition, these three drugs showed potential effects against a broad spectrum of pathogens in filamentous fungal keratitis, including Aspergillus [38], Purpureocillium [37], Beauveria [37], and Scedosporium [39] species. Moreover, because of their low molecular weight and lipophilicity, efinaconazole (348.39 Da), lanoconazole (319.8 Da), and luliconazole (354 Da) have theoretically good penetration into the corneal stroma and anterior chamber. However, currently, none of these new azoles are available as eyedrops. Only luliconazole is clinically available, in the forms of 1% ointment and 1% solution, for treating onychomycosis and dermatophytosis [40]. With low MICs against Fusarium species and theoretically good tissue penetration, these new azoles may be favorable candidates for filamentous fungal keratitis treatment in the future.

This study has several limitations. First, the clinical data might be incomplete because of the retrospective study design, and the physicians used diverse protocols in treating patients. Second, in vitro antifungal susceptibility may not be an accurate reflection of the clinical responses to keratitis because the concentration of antifungal agents may remain above the MIC on the ocular surface in cases of high-frequency topical use.

Third, although we incorporated *TEF-1* α as suggested by previous studies for the identification of these clinical *Fusarium* species [41,42], we found that *TEF-1* α alone failed to resolve some closely related species into well-supported monophyletic clades (e.g., *F. brevis* and *F. vanettenii*). In addition, some strains could not be classified as any known species in the ITS + *TEF-1* α analysis (e.g., CGMHD2044). Further investigation is required to determine whether these isolates are new *Fusarium* species. Moreover, our sample size was small, especially the non-FSSC group. We plan to include more cases to compare the clinical outcomes of FSSC and non-FSSC cases with greater statistical power in the future. Finally, as with other microbiological studies, our results may not be generalizable to other populations or territories because our study was conducted only at a single referral center in Taiwan. Nevertheless, this study is the first to provide an overview of *Fusarium* keratitis in

Taiwan, including molecular identification, antifungal susceptibilities, and clinical features. Moreover, only limited literature investigated the drug susceptibilities of efinaconazole, lanoconazole, and luliconazole against *Fusarium* keratitis isolates in vitro; our study not only provided supporting results for the potential of these new azoles in *Fusarium* keratitis treatment but also is the first to manifest the difference in their MICs between the FSSC and the non-FSSC isolates in vitro.

5. Conclusions

This epidemiological study analyzed *Fusarium* keratitis identified through molecular diagnostics at a referral center in Taiwan. The FSSC was the most common species complex, accounting for 76.7% of the *Fusarium* keratitis cases. In comparison with non-FSSC isolates, the MICs for FSSC isolates were higher for multiple antifungal agents, namely natamycin, voriconazole, chlorhexidine, luliconazole, and lanoconazole, but no significant differences in clinical outcomes were observed between FSSC and non-FSSC cases. Among the antifungal agents used in clinical practice, amphotericin B exhibited the lowest MIC, followed by natamycin and voriconazole. However, the three new azoles, efinaconazole, luliconazole, and lanoconazole, registered even lower MICs against *Fusarium* in vitro. Although these azoles are currently unavailable in eyedrop form, their broad-spectrum antifungal abilities and potentially good tissue penetration make them promising candidates for fungal keratitis treatment in the future.

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Appendix A. Appendix

		Case		Amphotericin B		Natamycin		Voriconazole		Chlorhexidine		Efinoconazole		Lanoconazole		Luliconazole	
		n.	%	Range	GM	Range	GM	Range	GM	Range	GM	Range	GM	Range	GM	Range	GM
FSSC	Total	33	76.7%	0.5~4	1.60	4~16	6.48	2~16	6.62	16~32	24.35	0.5~2	1.23	0.125~0.5	0.23	0.031~0.25	0.07
	F. falciforme	14	32.6%	1~2	1.35	8	8	2~16	5.22	16~32	27.27	0.5~2	0.90	0.125~0.5	0.17	0.031~0.25	0.05
	F. keratoplasticum	12	27.9%	2~4	2.12	4	4	2~16	6.35	16~32	22.63	0.5~2	1.50	0.125~0.5	0.24	0.031~0.125	0.07
	F. solani.	3	7.0%	0.5~2	1.59	4~16	11.31	8~16	13.45	16~32	22.63	1~2	2.00	0.25~0.5	0.42	0.063~0.25	0.15
	F. suttonianum	2	4.7%	1~2	1.41	8~16	11.31	16	16	16~32	22.63	2	2	0.5	0.5	0.063~0.125	0.09
	F. silvicola	1	2.3%	0.5	0.5	4	4	8	8	32	32	1	1	0.5	0.5	0.125	0.125
	F. breve	1	2.3%	2	2	8	8	2	2	16	16	0.5	0.5	0.125	0.125	0.031	0.031
FDSC	Total	4	9.3%	1	1	2~4	2.38	2	2	8	8	0.25~0.5	0.30	0.031~0.063	0.04	0.016	0.016
	Fusarium sp. FDSC ^a	3	7.0%	1	1	2	2	2	2	8	8	0.25	0.25	0.031~0.063	0.04	0.016	0.016
	F. delphinoides	1	2.3%	1	1	4	4	2	2	8	8	0.5	0.5	0.063	0.063	0.016	0.016
FOSC	Total	2	4.7%	2	2	4~8	5.66	2~8	4	8~16	11.31	0.25~4	1	0.063~0.25	0.13	0.031~0.063	0.04
	F. nirenbergiae	1	2.3%	2	2	8	8.00	8	8	8	8.00	4	4	0.25	0.25	0.063	0.063
	F. veterinarium	1	2.3%	2	2	4	4.00	2	2	16	16.00	0.25	0.25	0.063	0.063	0.031	0.031
FIESC	Total	2	4.7%														
	<i>Fusarium</i> sp. FIESC ^b	1	2.3%	1	1	4	4	4	4	8	8	1	1	0.25	0.25	0.063	0.063
	F. lacertarum	1	2.3%	No MIC	C data ^c												
FFSC	Total	2	4.7%	2	2	4~8	5.66	2~8	4	8~16	11.31	0.25~0.5	0.35	0.125	0.125	0.031~0.063	0.044
	F. aff. circinatum	1	2.3%	2	2	4	4	2	2	16	16	0.25	0.25	0.125	0.125	0.031	0.031
	F. annulatum	1	2.3%	2	2	8	8	8	8	8	8	0.5	0.5	0.125	0.125	0.063	0.063

Table A1. Antifungal susceptibilities for *Fusarium* keratitis isolates in vitro.

FSSC: *Fusarium solani* species complex; FOSC: *Fusarium oxysporum* species complex; FDSC: *Fusarium dimerum* species complex; FIESC: *Fusarium incarnatum-equiseti* species complex; FFSC: *Fusarium fujikuroi* species complex. ^a Hans-Josef Schroers et al., had identified this *Fusarium* species in 2009 but this species was still unnamed [24]. ^b The *TEF-1*α similarity of this isolate was only 96.3% to an isolate of *F. incarnatum* in Genbank (MAFF 236386), but in another article, the same species was renamed as *F. semitectum* [25]. Hence, further species identification was required for this isolate.

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