

V. CONCLUSIONES

Objetivo I

1. No ha sido fácil la recopilación de cepas de *Cryptococcus neoformans* ya que la mayoría de los laboratorios no las conservan. A pesar de todo se han obtenido 128 cepas de 115 pacientes, de 8 regiones geográficas de España. Varios de estos pacientes tenían más de un aislamiento en el tiempo. El período de los aislamientos abarca 10 años (1988-1997). El 95% de los aislamientos provienen de enfermos de sida.
2. La dificultad en la obtención de cepas de origen animal ha sido mayor. Se han obtenido 13 cepas aisladas de 6 cabras con criptococosis de 5 rebaños de diferentes áreas de la provincia de Cáceres.

Objetivo II

3. El estudio ambiental de muestras de polvo doméstico de pacientes con criptococosis, enfermos VIH+ y controles sanos ha resultado negativo. Este substrato no representa un reservorio saprofítico de *C. neoformans* en nuestro medio ambiente.

Objetivo III

4. La aplicación del medio de L-canavanina-glicina-azul de bromotimol junto con la asimilación de aminoácidos ha permitido demostrar por primera vez que la variedad *gattii* de *C. neoformans* existe en nuestro país. Esta variedad ha resultado patógena para un número elevado de animales produciendo brotes epidémicos de criptococosis localizadas en Cáceres. Todas las cepas estudiadas pertenecían al serotipo B.
5. El hallazgo de *C. neoformans* var *gattii* autóctono da soporte a que esta variedad no está restringida a zonas tropicales y subtropicales.

Objetivo IV

6. Se han obtenido anticuerpos policlonales de buena calidad en conejos para los serotipos A, B, C, y D con un costo muy inferior a los comercializados.
7. La distribución de los serotipos no es homogénea en España. El serotipo A es predominante en muestras clínicas y ambientales, no obstante se ha encontrado una elevada tasa del serotipo D principalmente en Valencia y Madrid.
8. No se han hallado hasta ahora los serotipos B y C en muestras clínicas humanas. En los aislados de cabras todas las cepas fueron del serotipo B.

Objetivo V

9. El nivel global de resistencias *in vitro* de las cepas clínicas ha resultado muy bajo. No se ha encontrado ninguna cepa con CIM > 1 µg/ml, para anfotericina B, sin embargo las del serotipo A mostraron menor sensibilidad a este polieno.
10. La 5-fluorocitosina ha resultado el antifúngico menos activo sobre *Cryptococcus neoformans*.
11. El fluconazol ha mostrado menor actividad *in vitro* que el itraconazol. Tres cepas mostraron CIM>64 µg/ml. El serotipo D muestra menor sensibilidad a los dos triazoles que el resto de los serotipos. ($P<0.00001$)
12. Las cepas de *C. neoformans* var. *gattii* aisladas de cabras muestran una sensibilidad uniformemente baja para los 4 antifúngicos estudiados.
13. Las cepas ambientales mostraron sensibilidades similares a las clínicas.

Objetivo VI

14. El método colorimétrico *Sensititre* es fácil en su ejecución y en la lectura de los puntos de corte. Este sistema proporciona valores más bajos de CIMs que el método de referencia.

Objetivo VII

15. Solamente 5 cepas de *C. neoformans* de las 190 aisladas en España y Latinoamérica mostraron CIM de 2 µg/ml para el nuevo antifúngico derivado del sordarín GM 237354. La variedad *gattii* mostró CIM, proporcionalmente más alta que los otros serotipos para el sordarín. Nuevamente no se hallaron valores ≥ 1 µg/ml. para la anfotericina B.

Objetivo VIII

16. La aplicación de la técnica de biología molecular del RAPD ha permitido demostrar que por lo menos existen 2 genotipos diferentes en la var. *gattii* aislada de cabras. Uno de los patrones parece ser el dominante ya que se ha hallado en todos los brotes epidémicos mientras que el otro patrón sólo se detectó en una zona.
17. La aplicación del RAPD en 7 cepas de 3 pacientes mostró su utilidad para discriminar entre diferentes aislamientos de un mismo paciente.

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Estudio de muestras de polvo doméstico para el aislamiento ambiental de *Cryptococcus neoformans*

Sr. Director,

La criptococosis es una infección fúngica que se adquiere habitualmente por vía inhalatoria. Esta infección se produce en la mayoría de los casos en personas inmunodeprimidas, en especial los pacientes con sida.

Cryptococcus neoformans presenta dos variedades, *C. neoformans* var. *neoformans*, agente causal de la criptococosis en humanos en nuestro país, cuyo nicho ecológico es con frecuencia las heces de aves sobre todo de paloma (*Columba livia*) [1] y *C. neoformans* var. *gattii*, que se ha relacionado con la presencia de *Eucalyptus* de diferentes especies, *Eucalyptus blakelyi*, *Eucalyptus camaldulensis*, *Eucalyptus gomphocephala*, *Eucalyptus rudis* y *Eucalyptus tereticornis* [2-4]. Hasta ahora no se ha descrito ningún caso de criptococosis en humanos por esta última variedad en España aunque sí que se ha observado en animales [5].

En la búsqueda del reservorio de *C. neoformans* var. *neoformans* se han descrito aislamientos a partir del polvo doméstico en ciudades como Kinshasa (Congo)[6] y Río de Janeiro (Brasil)[7]. En ambas se ha comprobado una elevada tasa de aislamientos en los domicilios de pacientes con sida, considerándose como una potencial fuente de infección. Para conocer si esta levadura se encuentra en el ambiente doméstico de Barcelona, se ha realizado un estudio de 79 muestras de polvo casero.

Se recogieron las muestras manualmente en un recipiente estéril del dormitorio y sala de estar del domicilio. Se separaron las muestras en tres grupos: 62 obtenidas del domicilio de personas sanas, 11 de personas VIH positivas y 6 de pacientes afectados de sida con criptococosis en el momento de la recogida.

Las muestras se procesaron a partir de una suspensión de 0,05 g de polvo en 15 ml de solución salina fisiológica estéril con 0,3 g/l de cloranfenicol [7]. Se sembraron con un escobillón en dos placas de Petri, una que contenía un medio con extracto de semillas de *Guizotia abyssinica* y otra con el medio preparado con semillas de girasol (*Helianthus annuus*) [8]. Ambos medios permiten la observación de colonias pigmentadas de color marrón, debido a la actividad de la fenoloxidasas por parte de *C. neoformans*. Las placas permanecieron a temperatura ambiente durante una semana y con observación diaria de las mismas.

Como control de calidad, se procesó, siguiendo la misma metodología, una muestra de polvo doméstico a la que se habían añadido 10⁶ de una suspensión en solución salina de *C. neoformans* a una concentración de 3 x 10⁶ UFC/ml así como una muestra de polvo procedente de Río de Janeiro (Brasil) que contenía esta especie. Once muestras de polvo doméstico correspondientes a dos pacientes con criptococosis y el resto de pacientes VIH + fueron remitidas al laboratorio de Micología de la Fundación Oswaldo Cruz (Río de Janeiro, Brasil), para su procesamiento. De las 79 muestras de polvo analizadas 33 fueron positivas para levaduras. En ningún caso se aisló *C. neoformans*. *Cryptococcus laurentii* fue aislado en tres muestras y *Cryptococcus albidus* en una. Otras levaduras aisladas han sido *Candida famata* (2 muestras), *Candida zeylanoides* (4 muestras), *Candida lipolytica* (1 muestra), *Candida* sp. (13 muestras), *Rhodotorula* sp. (8 muestras), *Aureobasidium* sp. (1 muestra). Tampoco fue aislado *C. neoformans* en las muestras procesadas en Brasil.

C. neoformans var. *neoformans* no ha sido aislado en el polvo doméstico de Barcelona, por lo que es de suponer que éste no constituye un reservorio de importancia para la transmisión de la criptococosis en ese medio. Posiblemente las condiciones bioclimáticas no favorecen el desarrollo de esta especie, al revés de 10 que sucede en regiones tropicales. Para determinar las fuentes naturales de la infección en España es necesario que se continúen realizando estudios ecológicos y epidemiológicos.

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First Identification of Autochthonous *Cryptococcus neoformans*
var. *gattii* Isolated from Goats with Predominantly Severe
Pulmonary Disease in Spain

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Cryptococcus neoformans var. *gattii* is associated with *Eucalyptus* trees growing in various tropical and subtropical regions of the world. The identification of 13 autochthonous strains of *C. neoformans* var. *gattii* in Spain is reported. These strains were isolated from lung (10 samples), liver (1 sample), and brain (2 samples) tissue specimens from six goats suffering from predominantly severe pulmonary disease that were autopsied. The animals were members of five different herds of goats grazing in rural areas of the province of Cáceres (Extremadura, Spain). Between 1990 and 1994, there were five outbreaks, in which between 2.5 and 12% of the goats were affected. Although respiratory symptoms (pneumonia) associated with cachexia were the predominant clinical picture in all outbreaks, brain and liver involvement was also documented in three of the five outbreaks. Biotyping was performed by culturing the isolates on L-canavanine-glycine-bromothymol blue medium and testing them for the assimilation of D-proline and D-tryptophan. Serotyping by agglutination tests confirmed the characterization of all strains as *C. neoformans* var. *gattii* serotype B. This is the first confirmation of the presence of this variety in Spain, with a peculiar ability to produce severe pulmonary and systemic disease in normal goats, particularly in the form of outbreaks of pneumonia in association with cachexia.

Cryptococcus neoformans is a capsulated yeast with a worldwide distribution. Since its description at the end of the last century, it has been isolated from different substrata from nature and it has been shown that the basidiomycete *Filobasi-diella neoformans* is its teleomorph (15). Humans and other animals are infected by inhalation, developing cryptococcosis, which is especially severe in immunocompromised individuals, in particular in those infected with the human immunodeficiency virus (4).

Two varieties of *C. neoformans* have been described, *C. neoformans* var. *neoformans* and *C. neoformans* var. *gattii* (6). Each variety has its distinctive serotypes based on the antigenic composition of its capsular polysaccharides, which play an important role in pathogenicity. *C. neoformans* var. *neoformans* consists of serotypes A, D, and AD, whereas *C. neoformans* var. *gattii* has the B and C serotypes (12). Substantial differences in the ecology of the two varieties have been described and noted to influence the epidemiology of cryptococcosis (17).

Cryptococcus neoformans var. *neoformans* has a worldwide distribution. It is isolated frequently from the droppings of birds, mainly the fecal pigeon (*Columba livia*), since it can grow in substrates containing high concentrations of creatinine (28). This variety is responsible for the majority of the cases of cryptococcosis in immunocompromised patients (19). By contrast, *C. neoformans* var. *gattii* has not been isolated from bird droppings, apparently because it has a lower tolerance for high

levels of creatinine. Its optimum growth temperature is 32°C, and plant debris are its natural reservoir, especially those of *Eucalyptus camaldulensis* (9) and *Eucalyptus tereticomis* trees (25). More rarely, it has been isolated from bat feces, a wasp's nest, and other substrata (10, 18). The epidemiology of the infections produced by *C. neoformans* var. *gattii* is also different, with infections appearing to occur predominantly in tropical and subtropical areas of Australia, Brazil, Kenya, Zaire, Southeastern Asia, and Southern California, affecting people with no impairment of their immunological status (8).

Cryptococcosis in wild and domestic mammals, with sporadic cases in cats, dogs, goats, horses, and sheep, has also been reported (3). Outbreaks in bovine and caprine livestock have also been identified (1, 23). Information regarding the variety of *C. neoformans* that causes infection in animals is scarce (11). In Europe, isolation of *C. neoformans* var. *gattii* is exceptional (20, 21).

We describe the first identification of *C. neoformans* var. *gattii* in Spain. The organism was isolated from lung, liver, and brain tissue specimens of six infected goats from five different herds in which five outbreaks of severe pulmonary and systemic disease occurred from 1990 to 1994.

MATERIALS AND METHODS

Organisms. Thirteen strains of *C. neoformans* isolated from tissue samples (obtained by autopsy) of six goats with subacute or chronic pneumonia associated with cachexia from farms in different districts were studied. These animals, of different *Capra hircus* races (*verala*, *selta/la*, and *murcia/la*) of local Spanish stock, were members of five different herds which had suffered from cyclic outbreaks of the disease between 1990 and 1994 in Cáceres, Extremadura, Spain. After their isolation and identification, the strains were kept in the collection of the Department of Infectious Diseases of the Faculty of Veterinary Medicine at the University of Extremadura for further studies.

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TABLE I. Characteristic of the different outbreaks of predominantly severe pulmonary disease associated with cachexia^a

Date of outbreak	No. of animals (%) ^b	Clinical prevalence per herd (%) ^c	Number of necropsies (m. of animal...)	Localization in central nervous system	Isolation of <i>C. neoformans</i> (%) ^d
1990	140	12	2	No	5
1991	250	2	1	Ycs	2
1994	3 (1)	10		No	2
1994	2 (1)	?		Ycs	2
1994	120	2.5	1	Ycs	2

^a The outbreaks in goats in different geographical areas of the province (n Cáceres, Extremadura, Spain.

^b Lethality rate. I(I)%.

^c Number of strains from different tissues.

firmed by histopathologic examination and culture. With the use of Indian ink, it was possible to observe encapsulated yeast cells characteristic of *C. neoformans* in pulmonary and brain tissue specimens. Thirteen very mucinoid strains were isolated from the 10 lung tissue samples as well as from the brain and liver tissues of the six autopsied goats. The identification showed that the 13 strains were *C. neoformans*, since the patterns of assimilation of sugars, cycloheximide tolerance, production of urease, and growth at 37°C were characteristic of this species. Strains were stained brown in the *H. annus* and *G. abyssinica* media. All of them were positive by the CGB test, and they assimilated D-proline and D-tryptophan, indicating that they belonged to *C. neoformans* var. *gattii*. Agglutination tests with the Crypto Check system demonstrated that all of the strains were serotype B. These results were confirmed by the Tropical Medicine Institute of Antwerp, Belgium.

DISCUSSION

Since Ellis and Pfeiffer (8, 9) isolated *C. neoformans* var. *gattii* from the environment in Australia, its natural habitat has been associated with *E. camaldulensis* (river red gum tree). This native species of eucalyptus is widespread in the south of Australia, and it has been exported and planted in areas of California, Mexico, and other parts of the world (24). The

isolation of *C. neoformans* var. *gattii* from nature has been reported from the tropical zones (of central Africa and Brazil (18). However, it has also been found in temperate zones of countries, such as Uruguay (10), and has been shown to cause some cases (of human cryptococcosis in Argentina (2). It has also been isolated from material obtained from another *Eucalyptus* tree, *E. tereticornis* and from different substrata, including a wasp's nest (10).

In Europe, *C. neoformans* var. *gattii* has been isolated in Germany from a patient working with wood of imported tropical trees (14); from environmental samples in Apulia, a region of Italy (21); and from *Eucalyptus* sp. samples in Portugal (20). Up to the present time, there have been no reports of the isolation of *C. neoformans* var. *gattii* from the environment or autochthonous infections in humans or other animals in Spain, a country with a temperate climate. *Eucalyptus* trees coming from France had been introduced in Extremadura since the last century, but an extensive reforestation with *Eucalyptus* sp. took place between 1955 and 1977. *E. camaldulensis* is by far the predominant species, followed by *Eucalyptus globulus* (5). Since 1980, no more reforestation campaigns with *Eucalyptus* have been promoted. Whether *C. neoformans* var. *gattii* was brought into Spain by infected imported trees is unknown.

It should be noted that autochthonous strains of *C. neoformans* var. *gattii* were isolated from goats of local Spanish stock suffering from severe subacute or chronic respiratory symptoms associated in some cases with systemic disease. The various animals from which this variety has been isolated were all found to be grazing free in different zones of the countryside in a region located in the central western part of the Iberian peninsula. The five outbreaks occurred in different geographical zones from 1990 to 1994. This suggests that there is wide distribution of this pathogenic yeast in the geographical area to which reference has been made. Inhalation was the most likely mode of infection. The animals in one of the herds had been grazing in a eucalyptus grove, in which the predominant species was *E. camaldulensis*.

In all cases, serotype B was identified. The prevalence of serotype B over serotype C in *C. neoformans* var. *gattii* everywhere except in the south of California is well known (9, 15, 17). Exhaustive studies to isolate this variety from the environment have still not been conducted in Spain, but undoubtedly this will become a primary objective in the near future.

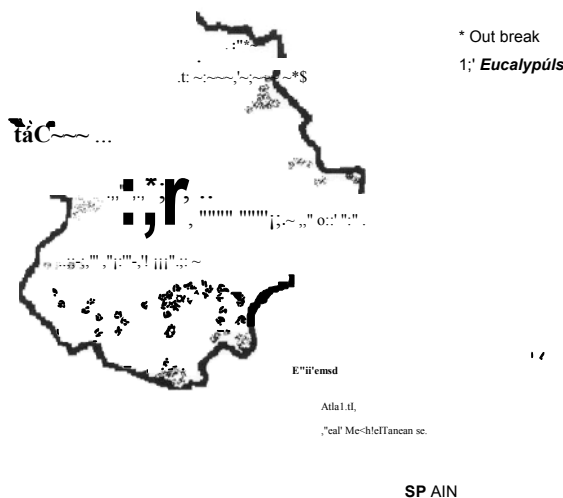


FIG. 2. Map of Extremadura, Spain, showing the locations of the five outbreaks of cryptococcal infections in goats and reforested areas with *Eucalyptus* spp.

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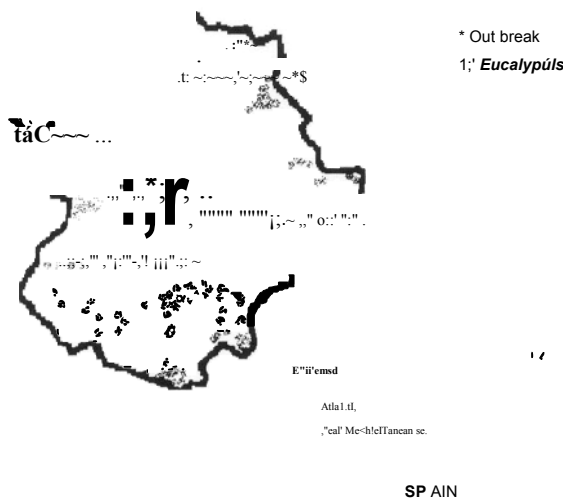


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TABLE 1. Oistribution of the scrutypes of 154 *C. neofonnans* isolates from Spain with respect to their sources

Source	Total no. of strains	No. (O/c) of strains of indicated serotype				
		A	D	AD	B	C
Humans	115		33 (29)		0	
Animals	6		0		6 (100)	0
Environment	33		5 (15)			

reaction for phenoloxidase in the niger seed and Pal's media. There was a complete agreement among the three methods used for determining the biovariety, and the biovars correlate with the serotypes. The strains of *C. neofonnans* var. *gattii* grew, producing a color change in CGB medium, and all as- simulated D-proline and D-tryptophan with enhanced growth around the disk. The serotype distribution of the Spanish iso- lates of *C. neofonnans* according to their source is shown in Table 1. Most of the human and environmental isolates be- longed to serotype A; only those isolated from goats were serotype B. Complete agreement in results was found with the 108 isolates tested with the Crypto Check agglutination meth- od. The geographical origin and serotype distribution of the clinical Spanish isolates are shown in Table 2. Distribution of clinical serotypes around the country was uneven (Fig. 1). Complete clinical data were obtained in 73% of the Spanish cases. Only three of them were HIV negative, and all belonged to serotype A. In the HIV -positive patients, serotype A was predominant (59.3%) but a high number of serotype D (28.4%) and serotype AD (12.3%) isolates were also found. No C serotype isolates were observed. Serotypes of the clinical isolates of *C. neofonnans* from Brazil, Cuba, and Argentina are shown in Table 3. *C. neofonnans* var. *gattii* was not found among the clinical isolates from Argentina and Cuba. One strain of *C. neofonnans* var. *gattii*, however, was obtained from a necropsied cheetah (*Acinonyx jubatus*) from the National Zoo of Havana. In contrast, 18 isolates of *C. neofonnans* var. *gattii* were identified among the Brazilian strains, 14 of them having been isolated from HIV -negative patients.

This was the first epidemiological survey performed in Spain on the prevalence and geographical distribution of the biova- rieties and serotypes of *C. neofonnans* .As in other European countries in which epidemiologic studies of *C. neofonnans* have been carried out, *C. neofonnans* var. *neofonnans* is the cause of cryptococcosis in HIV -infected patients and other immuno- compromised hosts. Until now, in Spain, no isolates of *C. neo- fonnans* var. *gattii* have been observed in humans, but this variety has been isolated from goats with cryptococcosis suf- fering from severe pulmonary disease (1). This finding suggests

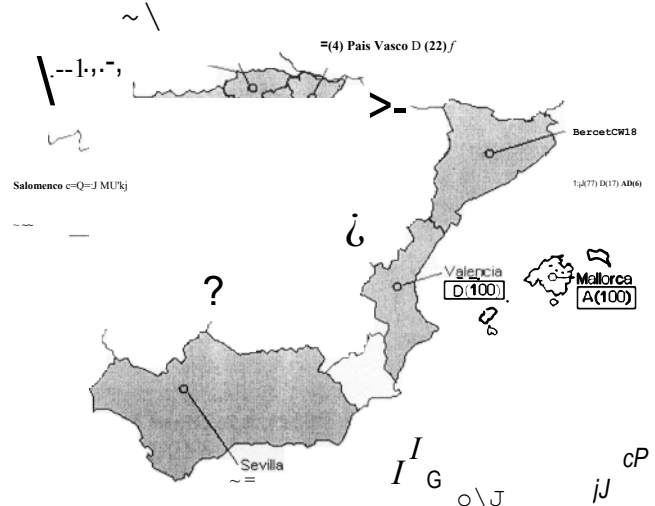


FIG. I. Oistribution of *C. neofonnans* clinical serotypes in Spain (results shown in percentages). Only single cultures from Santander and Salamanca were evaluated.

that the autochthonous distribution of serotype B of *C. neo- formans* is not limited to tropical and subtropical areas but also includes areas with temperate climates, such as Spain, Portu- gal, and Italy (23, 24). There seems to be a relationship be- tween the distribution of five species of *Eucalyptus*, including *E. blakelyi*, *E. camaldulensis*, *E. gomphocephala*, *E. rndis*, and *E. tereticomis* (11,27,28) and *C. neofonnans* var. *gattii* isolates. Most of these *Eucalyptus* species are found in some Spanish geographical locations, but no isolates of *C. neofonnans* var. *gattii* have been previously identified (6). In Europe, there have been a few reports in which autochthonous human isolates of *C. neofonnans* var. *gattii* have been described (18, 25). The three methods used for the differentiation of the two varieties were accurate, and the results were in agreement. All isolates of the *C. neofonnans* var. *gattii* assimilated D-proline and D- tryptophan and grew in CGB medium; therefore, any one of these methods could have been used. The assimilation test for D-amino acids is easy to perform and less expensive than growth in CGB medium.

Serotype A is prevalent in clinical specimens in Spain, but serotypes D (29%) and AD (9%) are frequent and predomi- nant in some areas (Fig. 1). This pattern also occurs with the environmental isolates, with serotypes D and AD frequently occurring. These data are consistent with the relative preva- lence of serotype D in some areas of France and Italy (8, 15, 33). We found an irregular distribution of serotype D among clinical isolates, with the highest prevalence of serotype D occurring in Valencia (eastern Spain) and Madrid (central

T ABLE 2. Geographical origin and serotype distribution of *C. neofonnans* among Spanish clinical isolates

Geographic region	Total no. of strains	No. (%) of strains of indicated serotype				
		A	D	AD	B	C
Barcelona	35	27 (77)	6(17)			
Valencia	11	0	11 (100)	0		
Mallorca	21	21 (100)	4 (20)	9		
Sevilla	20	12 (60)	(53)			
Madrid	17	6 (35)	1 (100)	2		
Salamanca	1	0	(22)			
Pais Vasco	9	4 (45)	0			
Santander	1	1 (100)				

TABLE 3. Distribution of *C. neofonnans* serotypes in human and animal isolates from a West Indian and two South American countries

Country	Total no. of strains	No. (%) of strains of indicated serotype				
		A	D	AD	B	C
Brazil	44	26 (59)				18 (41)
Cuba	22	21 (95)				1"
Argentina	44	39 (88)				(5) 0 0

u Isolats of animal origin.

Spain). No serotype C isolates were found either in Spain or Europe. Data from American isolates also showed a higher prevalence of serotype A than serotype D in Brazil and Cuba. The exception to this pattern was for isolates from Argentina, possibly because Buenos Aires has a temperate climate similar to that of Spain. We would like to draw attention to the first isolation of *C. neoformans* var. *gattii* in a cheetah from the National Zoo in Havana (Cuba). This animal is another mammalian victim of cryptococcosis.

Currently, studies using molecular biology techniques are in progress to establish the genomic DNA patterns of *C. neoformans* and thus obtain a better understanding of the epidemiology of cryptococcosis in Spain.

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RELATIONSHIP BETWEEN *IN VITRO* ACTIVITY OF FOUR ANTIFUNGAL DRUGS AND THE SEROTYPES OF *CRYPTOCOCCUS NEOFORMANS*

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SUMMARY ~

Objectives. The aim of this study was to assess the *in vitro* activity of four antifungal agents against 128 clinical strains of *Cryptococcus neoformans* isolated from different geographical regions of Spain. The susceptibility of strains has been related to the serotype found for each isolate.

Materials and methods. The MICs were determined by the broth microdilution technique by following the modified guidelines described in the National Committee for Clinical Standards (NCCLS) document M27-A, using RPMI medium + 2% glucose.

Results. There was no resistance to amphotericin B but serotype A was less susceptible to this antifungal agent ($P < 0.05$). Only 39% of isolates were fully susceptible to flucytosine, most of isolates being intermediate (60%). No statistical differences were found among serotypes.

The MIC in which 90% of the strains were inhibited by triazoles was 16 μ g/ml for fluconazole and 0.25 μ g/ml for itraconazole. Twenty-two percent of the strains had MICs \sim 16 μ g/ml (2% MIC \sim 64 μ g/ml) for fluconazole and 13% of the isolates had a MIC \sim 0.25 μ g/ml (0.9% MIC = 1 μ g/ml) for itraconazole. Serotype D was found to be less susceptible to both azoles ($P < 0.00001$).

Conclusion. The present study showed that the serotype A isolates of *C. neoformans* were less susceptible to amphotericin B than the serotypes D and AD ($P < 0.05$). There was no statistical significance in the susceptibility to flucytosine for the different serotypes. Most of *Cryptococcus neoformans* were not susceptible to this agent. For the triazole drugs, the serotype D is the least susceptible ($P < 0.00001$).

These results suggest a relationship between susceptibilities to some antifungal drugs and *C. neoformans* serotypes.

RÉSUMÉ : Relation entre l'activité *in vitro* de quatre antifongiques et le sérotype de divers isolats de *Cryptococcus neoformans*

Objectif. Le but de cette étude est de connaître l'activité *in vitro* de l'itraconazole. Vingt-deux pour cent des souches ont une CMI \geq 16 quatre agents antifongiques sur 128 souches cliniques de *Cryptococcus* μ g/ml (2% CMI \geq 64 μ g/ml) pour le fluconazole et 13% ont une *neoformans* isolées de différentes régions géographiques de l'Espagne, CMI \geq 0,25 μ g/ml (0,9% CMI = 1 μ g/ml) pour l'itraconazole. La sensibilité de chaque souche est comparée avec le sérotype trouvé, sérotype D montre la plus basse sensibilité pour les azolés

Matériels et méthodes. Les CMI sont déterminées par la technique ($P < 0.00001$),

méthode de microdilution en bouillon du "National Committee for Clinical Standards (NCCLS) document M27-A utilisant le milieu A de *C. neoformans* sont moins sensibles à l'amphotéricine B que les RPMI + 2% glucose. souches de sérotype D et AD ($P < 0.05$). Il n'y a pas de corrélation

Résultats. Il n'y a pas de résistance à l'amphotéricine B mais la statistique entre les différents sérotypes et la sensibilité à la flucytosine. sérotype A est moins sensible à cet antifongique ($P < 0.05$). Seulement la plupart des souches de *C. neoformans* ne sont pas sensibles à cet 39 % des isolats sont sensibles à flucytosine, la plupart des souches agent. Le sérotype D montre la plus faible sensibilité aux azolés (60 %) ont une sensibilité considérée comme intermédiaire. On n'a pas ($P < 0.00001$). trouvé de différence statistique entre les sérotypes pour cet agent Ces résultats suggèrent une relation entre la sensibilité *in vitro* à un antifongique. quelques antifongiques et les sérotypes de *C. neoformans*.

Par rapport aux triazoles, la CMI pour laquelle 90 % des souches sont inhibées est de 16 μ g/ml pour le fluconazole et de 0.25 μ g/ml pour

INTRODUCTION

Cryptococcus neoformans is an opportunistic fungal pathogen with a wide geographical distribution. This organism is the causative agent of cryptococcosis. Cryptococcal meningitis is the disease most frequently observed, but disseminated disease affecting any organ is also common, especially in human

immunodeficiency virus-infected patients (8, 21, 23). Primary treatment of acute disease with amphotericin B, with or without flucytosine, is usually recommended, however it is associated with a high toxicity profile and failure in some patients (10,34). To prevent relapses, long-term follow-up prophylaxis mainly with fluconazole (17) is used.

The development of standard methods for routine susceptibility testing of *C. neoformans* is important to facilitate the choice of antifungal agents and to select the appropriate doses of these agents, particularly in the case of resistant isolates (9, 32).

Few studies on the antifungal susceptibilities of clinical strains of *C. neoformans* have been performed in Spain (30) and in any case there is no information on the serotype distribution

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of isolates. The aim of the present work is to assess the *in vitro* activity of four antifungal agents, (amphotericin B, flucytosine, fluconazole and itraconazole) against 128 clinical strains of *C. neoformans* isolated from different geographical regions of Spain by using the reference microdilution method proposed by the National Committee for Clinical Laboratory Standards (NCCLS, M27-A document) (25). The susceptibility of each strain has been related to the serotype found for each isolate.

MATERIALS AND METHODS

Antifungal agents

Amphotericin B was provided by Squibb (Esplugues LL., Spain). flucytosine by La-Roche (Hoffman-La Roche, Switzerland), fluconazole was from Pfizer Inc. (Central Research, Sandwich, U.K.), and itraconazole from Janssen (Research-Foundation, Belgium). Amphotericin B and itraconazole were prepared in 100% dimethyl sulfoxide (DMSO) and flucytosine and fluconazole in sterile water. Stock solutions were prepared as described in document M27-A (25). The antifungal agents were dissolved at a concentration 100 times that of the highest dose to be tested (1.6 mg/ml for amphotericin B and itraconazole; and 6.4 mg/ml for flucytosine and fluconazole) and stored at -70 °C. until thawing on the day of susceptibility testing.

Medium

RPMI 1640 medium (Sigma Chemical, St Louis, Missouri 63103 USA) with glutamine but not bicarbonate + 2% glucose buffered to pH 7 with 0.165 M/L N-morpholino propanesulphonic acid (MOPS) buffer (Sigma) was used in all studies.

Microtiter plates

Microbroth dilution testing was performed on sterile microdilution plates, (Nunc@J, Brand products, DK-4000 Roskilde-Denmark). Dilutions of each antifungal agent were made in RPMI 1640 medium and 100 μ l dispensed into wells 1 to 10 of each row. The antifungal concentrations ranged from 0.03 to 16 μ g/ml for amphotericin B and itraconazole, and 0.125 to 64 μ g/ml for flucytosine and fluconazole, and were obtained by 10 two-fold serial dilutions. Drug-free medium was dispensed into wells 11 and 12. Well 11 served as a sterility control and spectrophotometric blank and well 12 served as a growth control.

Patients and Organisms

One hundred and twenty-two clinical isolates of *C. neoformans* from different patients and six strains from three patients who had two different serotypes in successive episodes of cryptococcosis were tested to determine drug MICs. These isolates were obtained between 1988 and 1998 from different patients of 11 Spanish hospitals in Catalonia, the Basque Country, Valencia, Majorca, Andalusia and Madrid. In 95% of the cases, samples were obtained from human immunodeficiency virus (HIV) infected patients. All of them were intravenous drug users except for one haemophilic and one homosexual patient. The non-HIV-infected patients included recipients of organ transplantation as well as patients with haematological malignancies or solid tumours. The isolates had been previously identified according to standard procedures (5, 36). *Cryptococcus neoformans* (ATCC 90112), *Candida albicans* (ATCC 90028) and *Candida krusei* (ATCC 6258) were included as reference or quality control strains whenever a test was run.

Serotyping

The serotypes of *C. neoformans* were tested by a slide agglutination test with polyclonal sera prepared in our laboratory (6) and confirmed

with a Crypto Check agglutination test (Iatron Labs Inc., Tokyo, Japan) (18, 20).

Inocula

All the isolates and control strains were grown on Sabouraud dextrose agar. (bioMérieux SA, 69280 Marcy l'Étoile-France) at 35 °C. for 48 h. The yeast inocula were adjusted to the 0.5 McFarland standard (1×10^8 - 5×10^8 CFU/ml) confirming the cell density with the aid of a spectrophotometer. The yeast stock suspension was mixed for 15 seconds with a vortex mixer. and diluted 1: 50 and further diluted 1: 20 with RPMI 1640 medium to obtain the inoculum (1×10^7 to 5×10^7 CFU/ml). This inoculum was diluted 1: 1 when the wells were inoculated to obtain a concentration of 0.5×10^7 to 2.5×10^7 CFU/ml. An aliquot of 0.1 ml was added to each well of the microdilution tray.

Incubation times

The plates were incubated at 35 °C, and microdilution MICs were read after 48 h. for *C. albicans* and *C. krusei* and after 72 h. for *C. neoformans*.

Endpoint criteria

The readings of the endpoints were performed visually using a reading mirror and the absorbances were determined spectrophotometrically (A420) after automatic agitation of the plates for 15 seconds with an automatic microplate reader (Labsystems, Multiskan MS, Finland). The MIC for amphotericin B was the lowest drug concentration in which there was absence of turbidity. The MIC of flucytosine and azoles was defined as the lowest drug concentration in a well that produced 80% reduction in turbidity compared with growth control (25, 27, 35).

As the breakpoint susceptibility values for *C. neoformans* have not yet been proposed by the NCCLS we adopted the breakpoint values proposed by the NCCLS document M27 A for *Candida* spp. to *Crypto- coccus* (9, 26,37) in order to have a defined group of values for analysing the results.

All the isolates presenting unusually high MICs were tested twice for confirmation of results.

Statistical analysis

The chi-square test was used to compare proportions in contingency tables. Whenever an expected value was less than 5. Fisher's exact test was used.

Kruskal-Wallis non-parametric ANOVA or the Mann-Whitney "U" test were used to compare groups medians for continuous variables that did not fit a normal distribution.

RESULTS

The susceptibility of 128 clinical isolates of *C. neoformans* was tested with four antifungal drugs. *Table I* describe the MICs required to inhibit 50 and 90% of the isolates and the range of control strains for all antifungal drugs. MICs for the control strains agreed with the NCCLS proposed values (24).

The MIC distribution of amphotericin B, flucytosine, fluconazole and itraconazole are presented (*fig. 1*). Two isolates of *C. neoformans* serotype A from Andalusia showed the highest MICs (11 μ g/ml) to amphotericin B, the MIC₉₀ was 0.51 μ g/ml for this polyene. Flucytosine was less active against *C. neoformans* than amphotericin B and azoles (P < 0.0001). Only 39% of strains presented a MIC \sim 4 μ g/ml to flucytosine, 60% with an

TABLE I. -/1/ \.lirO aclivlie,f (iffi)ur a/llifu/lgal agens agai/1sl 128 Cl)plOCOCcu.f /leofOrman.f cli/1ical i.folale.f a/ld Ihree reference .flrain.f. Aclivilé in I.ltro de qualre anlfo/lgiqille.f .fUr 128 isola.f clinique.f de Cl)'PIOCOCCHIS ne(-f)'nan.f el sur I.rois .f'!lIChe.f de référence.

Agent and strain group		MIC (J/g/ml)*	90%
		Range	
Amphotericin B			
Clinical		0.03-1	0.125
<i>C. albicans</i>	ATCC 90112	0.25-0.5	0.5
<i>C. albicans</i>	ATCC 90028	0.5-1	
<i>C. krusei</i>	ATCC 6058	0.5-1	
Flucytosine			
Clinical		0.25-32	
<i>C. albicans</i>	ATCC 90112	2.8	
<i>C. krusei</i>	ATCC 90028	0.5-1	
Fluconazole	ATCC 6058	8-16	
Clinical		0.5- > 64	8
<i>C. albicans</i>	ATCC 90112	2-8	16
<i>C. krusei</i>	ATCC 90028	0.25-1 32-	
Itraconazole	ATCC 6058	64	
Clinical		0.03-1	0.06
<i>C. albicans</i>	ATCC 90112	0.03-0.06	0.25
<i>C. krusei</i>	ATCC 90028	0.03	
	ATCC 6058	0.12-0.25	

* 50% and 90% MICs for which 50 and 90% of the strains, respectively, are inhibited.

MIC of 8-16 µg/ml, and one clinical isolate gave an MIC of 32 µg/ml. This isolate was serotype D and came from Madrid. The study of susceptibility to the triazoles showed that fluconazole had MICs S 8 µg/ml for 78% of the *C. neoformans* isolates tested (100 of 128), 16 to 32 µg/ml for 20% of these isolates (26 of 128), and ?; 64 µg/ml for 2% (3 of 128). Two serotype A isolates and one serotype AD isolate had a MIC ?; 64 µg/ml to fluconazole. Itraconazole had MICs of S 0.12 µg/ml for 87% of the isolates (111 of 128), 0.25 to 0.5 µg/ml for 12% (16 of 128) and only one serotype A isolate which was from Catalonia (Barcelona) had a MIC of 1 µg/ml.

Of all geographical areas, the isolates from Valencia showed the highest susceptibility (73%; $p < 0.05$) to flucytosine. With respect to the azoles, none of these isolates showed a MIC S 0.125 µg/ml to itraconazole and a MIC S 8 µg/ml to fluconazole. Neither showed values of MIC ?; 1 µg/ml and ?; 64 µg/ml respectively ($P < 0.0001$).

The Kruskal-Wallis test was used for comparing the MICs of the antifungal agents with respect to the different serotypes. This analysis revealed that although all the isolates were sensitive to amphotericin B, serotype A tended to have a lower susceptibility (MIC median 0.25 µg/ml) than the other serotypes D and AD, (MIC median 0.125 µg/ml) with $p < 0.05$. In spite of the fact that some investigators such as Nguyen and Yu (26) have suggested MIC values for *C. neoformans* equivalent to those of *Candida* spp. in order to consider an isolate as susceptible, intermediate or resistant, in this study three break-

points to antifungal drugs were defined, for flucytosine. MIC ~ 4 I.Lg/ml. 8-16 I.Lg/ml and ?; 32 I.Lg/ml. for fluconazole MIC ~ 8 I.Lg/ml. 16-32 I.Lg/ml and ?; 64 I.Lg/ml. and for itraconazole MIC ~ 0.125 I.Lg/ml. 0.25-0.5 I.Lg/ml and ?; 1 I.Lg/ml. These breakpoints were compared with the serotypes by using a chi-square test (fig. 2). There were no significant differences among the serotypes and susceptibilities for flucytosine. In contrast, serotype D was less sensitive to the azoles ($P < 0.00001$) as 60% of the strains were in the interval 16-32 I.Lg/ml to fluconazole and 0.25-0.5 I.Lg/ml to itraconazole.

DISCUSSION

The *in vitro* susceptibilities of *C. neoformans* to amphotericin B, flucytosine, fluconazole and itraconazole have already been described by different investigators (4, 7, 38). However, only very limited information is available for some countries such as Spain. As the ecology of *C. neoformans* and the epidemiology of cryptococcosis are different in many parts of the world it is important to know whether such differences also reflect variations in susceptibilities to antifungal drugs.

One would expect the breakpoints selected for *Candida* not to be useful when applied to other organisms such as *C. neoformans*. However, some papers use this criterion to evaluate the susceptibilities of *C. neoformans* (9, 26). It is generally accepted that *in vitro* values can influence the effectiveness of

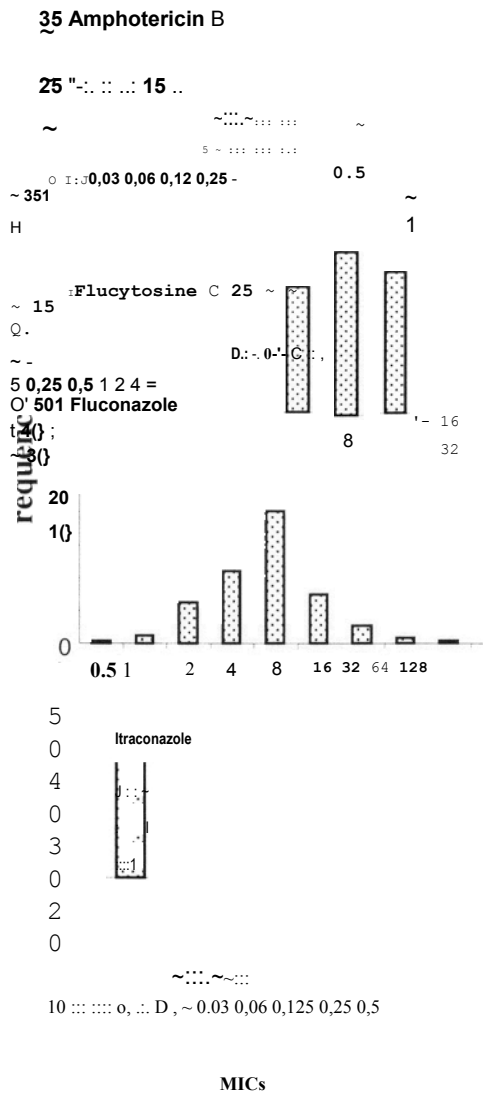


FIG. 1. -Distribution of MICs (11g/ml) of amphotericin B, flucytosine, fluconazole and itraconazole for 128 clinical isolates of *Cryptococcus neoformans*.

Distribution des CMIs (11g/ml) de l'amphotéricine B, de la flucytosine, du fluconazole et de l'itraconazole ou 128 isolats cliniques de *Cryptococcus neoformans*. susceptible strains do not always predict therapeutic success, but a high endpoint of *C. neoformans* will often predict therapeutic failure. (1, 13, 15, 19, 28,33, 39).

The relationships between serotypes of *C. neoformans* and antifungal susceptibilities are not well known. Poonwan *et al.* (31) studied the susceptibility of 50 selected strains of *C. neoformans* isolated from clinical specimens in Thailand and found only one resistant strain for flucytosine. but the authors did not indicate the serotype of this isolate.

The aim of the present study was to determine the MICs of the four most frequently used antifungal drugs for treating cryptococcosis. according to the different serotypes found in human clinical isolates of *Cryptococcus neoformans* in Spain. An additional objective was to relate the determined MICs to the uneven geographical distribution of cryptococci in this country (6). as also was described for Dromer *et al.* in France (II).

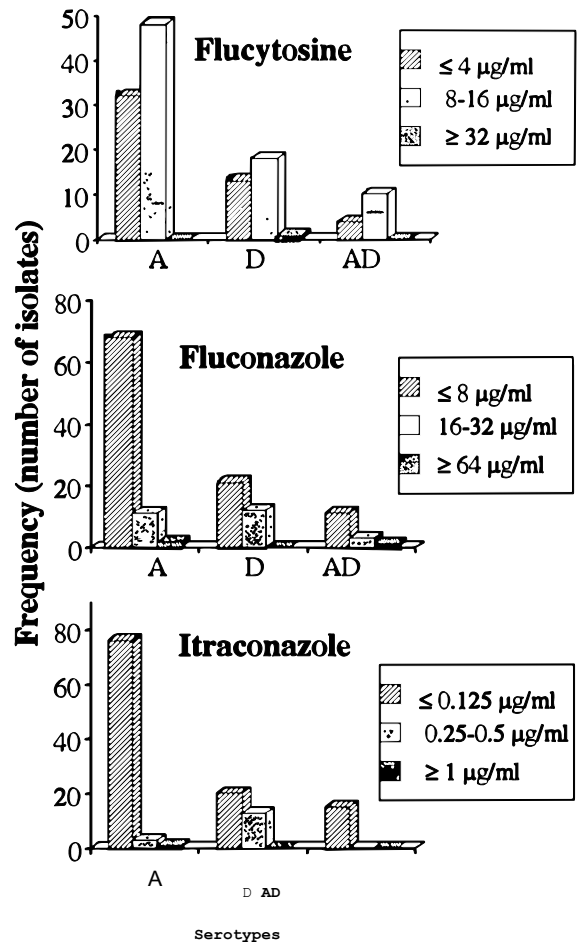


FIG. 2. -Susceptibility of the serotypes A, D and AD of *Cryptococcus neoformans* for flucytosine, fluconazole and itraconazole.

Sensibilité des sérotypes A, D et AD de *Cryptococcus neoformans* a la flucytosine, au fluconazole et a l'itraconazole.

Cryptococcus neoformans serotype B isolates were not found in humans, although they have previously been isolated in Spain from animals (6).

Susceptibility could be determined by using different methods but the standardised NCCLS M27-A method was chosen because comparisons with others studies can be made. and also because it offers the possibility of adapting the categories of susceptibilities for *Candida* spp to *C. neoformans* as other authors have published (9, 26). Obviously this categorisation of susceptibility must be confirmed by other studies including therapeutic responses (clinical aspects and treatment).

In agreement with other investigators (2, 14, 16) and based on our own experience, we found that RPMI 1640 medium with 2% glucose is a suitable medium for testing cryptococci. Although other authors suggest that Yeast Nitrogen Base (YNB) may enhance the growth of *C. neoformans*. in a pilot study we did not find any differences between RPMI and YNB (results not shown).

The results for the *in vitro* activity of amphotericin B showed that there were no *Cryptococcus*-resistant isolates to this poly- ene. Other authors have however found resistance to this drug

(9, 22, 32). Flucytosine gave the highest MICs compared with the other drugs (3), but only 0.8% of the isolates had MIC \geq 32 μ g/ml and 60% of the strains were in the MIC interval 8-16 μ g/ml.

Resistance of *C. neoformans* var. *neoformanis* has begun to emerge in patients undergoing prolonged azole treatment (28). In our study only three isolates presented MICs \geq 64 μ g/ml to fluconazole. If we compare these results with similar studies (29), it seems that the Spanish isolates were less sensitive to fluconazole (78%) than the African isolates (94%), and similar to the U.S. isolates (80%). The possibility of acquired resistance in patients on prolonged prophylactic fluconazole treatment should be considered. These results could be related to the extensive use of this azole in patients infected with human immunodeficiency virus (HIV) in Spain especially before the generalisation of the High Activity Antiretroviral Therapy (HAART). None of the patients included in this study ever received this kind of retroviral therapy.

In our study 95% of patients were HIV positive and we can assume that most of them had received antifungal treatment during the course of the disease, especially fluconazole for oropharyngeal candidosis. Considering that the samples were obtained over a period of 10 years from 11 hospitals from various geographical regions of Spain, there is no unified antifungal therapy regime, nor primary prophylaxis, rejected by the majority of Spanish clinics.

Itraconazole showed higher *in vitro* activity in our study than fluconazole, only one isolate was found with a MIC = 1 μ g/ml to this triazole. This strain was found to have a MIC of 32 μ g/ml to fluconazole. Itraconazole is rarely administered for the treatment of cryptococcosis in Spain.

Regarding the serotypes, there were no resistant isolates to amphotericin B using the present methodology, but the statistical analysis showed that the serotype A isolates of *C. neoformans* were less susceptible to this polyene than the serotype D and AD isolates ($P < 0.05$). So far this finding does not seem to be clinically relevant but could reflect a tendency in the follow up.

There was no statistical significance in the susceptibility to flucytosine for the different serotypes, and the only resistant isolate to this drug was from the serotype D.

Dromer *et al.* (12) suggest that individual and environmental factors may be associated with the serotype of *C. neoformans*. We found that, for the azole drugs, serotype D is the least susceptible ($P < 0.00001$). As the conditions of patients were similar (HIV + and the frequency of use of fluconazole for oropharyngeal candidosis) the MIC differences for azoles should be attributed to serotypes. These results however should be confirmed by a higher number of isolates.

The present results suggest that it is relevant to know the status of the susceptibility and resistance of *C. neoformans* to antifungal drugs in different parts of the world because significant differences could exist due to ecological and local factors as well as the antibiotic usage policy. Surveillance is also necessary for detecting the emerging resistance of *C. neoformans* strains to present or new antifungal drugs.

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***In vitro* susceptibility of *Cryptococcus neoformans* isolates to five antifungal drugs using a colorimetric system and the reference microbroth method**

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Minimum inhibitory concentrations (MICs) of amphotericin B, 5-flucytosine, fluconazole, itraconazole and ketoconazole were determined against 42 clinical isolates of *Cryptococcus neoformans* var. *neoformans* using the Alamar YeastOne colorimetric method and the NCCLS reference microdilution method. No strains with resistance to amphotericin B, itraconazole or ketoconazole were detected with either method. Using the reference method, the MICs of fluconazole were ~ 64 mg/L, whereas using the colorimetric method all MICs were ~ 16 mg/L. The MIC values of 5-flucytosine were also higher using the reference method (8-16 mg/L for 32% of isolates) compared with the colorimetric method. The percentage of agreement between the methods, using a difference of two dilutions, was 70.7% for itraconazole; 73.2% for amphotericin B, 80% for fluconazole, 88% for 5-flucytosine and 95% for ketoconazole. Overall, we conclude that for fluconazole and 5-flucytosine, in a low but not insignificant number of isolates, results with the two methods are discordant, some isolates being found sensitive with the colorimetric test, but resistant with the reference method.

Introduction

Strains of *Cryptococcus neoformans* resistant to antifungal agents including flucytosine, fluconazole and amphotericin B have been shown to account for failure or relapse during azole treatment of cryptococcosis. This makes antifungal susceptibility testing even more important in selecting and monitoring antifungal chemotherapy. Although standardized methods for broth macrodilution and microdilution testing of yeasts have been developed by the National Committee for Clinical Laboratory Standards (NCCLS), the availability of commercially prepared microdilution antifungal panels has been shown to enhance the ability to detect resistant isolates. Recently, a dried microdilution panel that incorporates an oxidation-reduction indicator (Alamar Bioscience Inc., Sacramento, CA, USA) has become available. Alamar YeastOne (Sensititre-Alamar YeastOne, Westlake, OH, USA) is a ready-to-use product; the 96-well microtitre plate has a colorimetric agent incorporated with the dried drugs. This produces a change from blue to pink in response to chemi-

cal reduction in the growth medium caused by the growing organisms in the inoculated wells. The first blue well in a row corresponds to the MIC of that drug. The antifungals included in the kit are amphotericin B, 5-flucytosine, fluconazole, itraconazole and ketoconazole. The Alamar colorimetric technique has demonstrated excellent agreement with the reference method when testing Gram-negative bacteria as well as for the susceptibility testing of yeasts and a high degree of intra- and interlaboratory reproducibility.

We evaluated the reliability of the results obtained by the Alamar colorimetric method compared with the results generated by the reference microdilution method for 42 clinical isolates of *C. neoformans*.

Materials and methods

Isolates

A total of 42 clinical isolates of *C. neoformans* var. *neoformans* (serotypes A, D and AD) obtained from AIDS

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patients during the initial diagnosis of cryptococcal infection were tested. Quality control was ensured by testing *Candida albicans* ATCC 90028, *Candida krusei* ATCC 6258, *Candida parapsilosis* ATCC 22019 and *Candida lusitanaensis* ATCC 90112 strains. Before testing, all isolates were subcultured on to Sabouraud dextrose agar (bio-Mérieux, Marcy l'Etoile, France) to ensure optimal growth characteristics. Stock suspensions were prepared in sterile phosphate-buffered saline (PBS) and adjusted to give a final inoculum concentration of 1×10^6 – 5×10^6 cells/mL.

Colorimetric method

The Alamar colorimetric method was performed according to the manufacturer's instructions. Working suspensions were prepared by adding 20 μ L of stock yeast suspension to 11 mL of RPMI 1640 broth (American Biorganics, Niagara Falls, NY, USA) buffered to pH 7.0 with 0.165 M morpholinepropanesulphonic acid (MOPS). This resulted in a final inoculum of 1.5×10^6 – 8×10^6 cells/mL. The final concentrations of the antifungal agents were 0.04–4 mg/L of amphotericin B, 0.12–256 mg/L of fluconazole, 0.04–64 mg/L of 5-flucytosine and 0.008–16 mg/L of itraconazole and ketoconazole. The wells were reconstituted by the addition of 100 μ L of the inoculum suspension. After incubation at 35°C for 48 h for *Candida* spp. and 72 h for *C. neoformans*, MICs were determined by observing the lowest antifungal concentration preventing the development of a red colour (first blue well).

Reference microbroth method

The microdilution method was performed according to the recommendations of M27-A 7 using RPMI 1640 medium and 2% glucose in MOPS buffered to pH 7. Dimethyl sulphoxide (DMSO) was used as solvent for the antifungals, the stock solutions were prepared at 100 X the highest concentration to be tested. The final concentration was prepared from the antifungal stock solution in RPMI plus 2% glucose. The antifungal agents were dispensed in sterile, individually wrapped polystyrene round bottom assay plates. The stock solutions of antifungal agents were dispersed in the assay medium to obtain appropriate concentrations in wells 1–10 in each row; drug-free medium was dispensed in wells 11 and 12. Well 12 served as sterility control and well 11 as growth control. The antifungal concentrations were 0.03–16 mg/L of amphotericin B (Squibb, Princeton, NJ, USA), ketoconazole and itraconazole (Janssen Biotech, Beerse, Belgium), 0.125–64 mg/L of fluconazole (Pfizer Inc, New York, USA) and 5-flucytosine (La-Roche Laboratory Inc., Nutley, NJ, USA). The yeast inoculum was adjusted to 0.5 McFarland standard. A working suspension was made at 1:100 dilution followed by a 1:20 dilution of a stock suspension with RPMI 1640 plus 2% glucose.

The inoculated plates were incubated for 72 h at 35°C and readings were taken daily. Absorbance was determined spectrophotometrically at 420 nm after agitation of the plates. The MIC endpoint was defined as the lowest drug concentration exhibiting approximately 80% (or more) reduction of growth compared with the control well. For amphotericin B the MIC was defined as the IOWC1 concentration giving 100% inhibition (optically clear).

Analysis of results

Discrepancies between MIC endpoints of no more than two dilutions were used to calculate the percentage agreement between the Alamar colorimetric method and the NCCLS reference microdilution method. The Student's *t* test for paired data was used for statistical analysis. Statistical significance was set at $p < 0.05$. The SPSS compute program was used for analysis of data.

Results

After 48 h incubation, MIC for *Candida* spp. were within the expected ranges in both the reference microdilution method and the Alamar colorimetric method. By the Alamar colorimetric method, 40% of *C. neoformans* isolate failed to produce any colour change in the growth control well after 48 h incubation, so readings were performed after 72 h incubation. Because the MIC scale is two dilutions (0.016 and 0.008 μ g/L) lower than that of the reference method, values \sim 0.03 mg/L were considered equivalent. Table I summarizes MIC ranges of the five antifungals tested against 42 *C. neoformans* isolates determined by the reference microdilution method and the Alamar colorimetric method. The results are reported as MIC ranges, MIC₅₀ and MIC₉₀, respectively. The highest MIC values were found for fluconazole and the lowest for ketoconazole. The levels of agreement between the two methods for *C. neoformans* isolates are given in Table II. Maximum and minimum agreements were found for ketoconazole (95.10%) and itraconazole (70.70%), respectively.

There were statistically significant differences between the methods in both the range and distribution of MICs for amphotericin B and 5-flucytosine, with a wider range and significantly higher mean MIC values for the microdilution reference method as compared with the Alamar colorimetric technique (amphotericin B, 0.143 versus 0.026 mg/L $p < 0.005$; 5-flucytosine, 6.316 versus 2.06 mg/L, $p < 0.0005$). In respect of the three azole compounds, lower mean MIC values ($P = 0.08$) were obtained for fluconazole with the Alamar colorimetric method than with the reference method, in which MIC values \sim 64 mg/L were obtained in 2.3% of strains. Differences between the three azole agents were greater for MIC₉₀ than MIC₅₀. All compounds showed identical MIC₅₀ values.

Table I. MIC distribution in 42 isolates of *Cryptococcus neoformans* against five antifungal drugs with the micro Alamar-Blue colorimetric test (data in percentages)

MIC (mg/L)	amphotericin B Alamar		fluconazole		itraconazole		Alamar NCCLS	
	NCCLS	Alamar	Alamar NCCLS	Alamar	Alamar NCCLS	Alamar		
0.008							7.1	
0.016	38				2.3		14.3	
0.03	57.1	31.11			9.5	57.14	2	
0.06	4.7	28.89	2.3		4	11.90	33.3	
0.125		13.33	2.3		5	14.29	3	
0.25		22.22			3	9.52	38	
0.5		4.44	2.3	4.26	3	7.14	4.7	
1			7.1	17.02	7.1		2.3	
2			14.2	10.64	2.3			
4			26.1	25.53				
8			38	23.				
16			4.7	4				
32				10.64				
~64				6.38				
Range Mean	0.016-0.06	<0.03-0.5	0.06-16	0.026 0.143 7.431	2.13	0.016-0.5	<0.03-0.5	0.08-0.2
MIC MIC50	0.03 0.06	4			0.5-64	0.101	0.10	0.047
MI~	0.03 0.25	8	0.03	0.06	8.44	0.06	0.06	0.03
					4	0.25	0.125	0.06
					1			
					6			

Table II. Agreement and differences (%) between the colorimetric and microdilution methods for five antifungal drugs: amphotericin B, fluconazole, itraconazole, voriconazole and 5-flucytosine against *C. neoformans* isolates

Dilutions		Amphotericin B	Fluconazole	Itraconazole	Voriconazole	5-Flucytosine
3	agreement	92.6%	92.5	100	100	95.4%
	difference	7.3	7.5	0	0	4.6%
2	agreement	73.171	80	70.732	95.12	11.57%
	difference	26.829	20	29.268	4.88	88.43%
	agreement	43.902	72.5	43.902	75.61	16.14%
	difference	56.098	27.5	56.098	24.39	83.86%
	agreement	17.073	22.5	9.7561	24.39	6.86%
	difference	82.927	77.5	90.244	75.61	93.14%

Discussion

There are only a few studies on the susceptibility of *C. neoformans*, so it is interesting to have data available on isolates from different sources and geographical distributions. To compare results, however, it is necessary to use a similar and standardized methodology. Although the reference methods provide objective and comparable results, the availability of automated plate-reading technology simplifies the procedures including MIC endpoint determination and avoids many manipulations of the standardized reference methods. In isolates of *Candida* spp., the recently introduced colorimetric micromethod has shown a good reproducibility and correlation with macrodilution and microdilution methods. In a study similar to ours, Pfaller & Barry [10] compared the NCCLS reference method and the Alamar colorimetric technique in 600 clinical yeast isolates. The collection, however, included only four *C. neoformans* isolates. These authors found a 100% agreement between both methods in MIC50 of amphotericin B, fluconazole and 5-flucytosine. It should be noted that all four *C. neoformans* isolates were susceptible to all antifungal agents with MIC ranges 0.25-1.0 mg/L.

Testing of the susceptibility of *C. neoformans* to different drugs has been extensively studied with a variety of culture media, although the number of published reports of clinical resistance to amphotericin B in *C. neoformans* is surprisingly low. Lozano-Chiu et al. have shown that both substitution of RPMI 1640 for antibiotic medium 3 in the microdilution variant of the M27-A method and use of the Etest agar diffusion methodology permits detection of amphotericin B-resistant *Candida* isolates. Only antibiotic medium 3, however, permitted consistent detection of amphotericin B-resistant *C. neoformans*. Because RPMI 1640 is the medium used in the colorimetric method, this may be the reason for the low number of resistant isolates found in the present study. Although all isolates were susceptible to amphotericin B, MIC ranges were lower for the Alamar test than for the reference microdilution method in which the RPMI 1640 medium was also used.

Similar findings were obtained with 5-flucytosine, MIC (varying between 0.06 and 4 mg/L for the Alamar test) at between 0.5 and 16 mg/L for the reference method (33% isolates showed MIC values ~8 mg/L). These differences were even greater for MIC50.

With amphotericin and itraconazole the distribution of MICs for both methods was largely different. It should be noted, however, that no discrepancies in resistance (sensitivity) of *C. neoformans* strains to these two antifungal agents were observed. In contrast, MIC values of fluconazole and 5-flucytosine obtained with the Alamar test and the reference method were quite similar, but with the important drawback that some strains apparently sensitive to fluconazole and 5-flucytosine with the Alamar test were resistant to the agents when tested with the reference method.

If the MIC breakpoints for fluconazole resistance in *C. albicans* mucosal infections (~64 mg/L) are applied, the number of *C. neoformans* strains classified as resistant was low. Davey et al. have reported an incidence of 5.6%, so that it may be expected that the 2.3% found in the present study with the reference method is more close to the actual situation than the total absence of resistance observed with the colorimetric micromethod.

The degree of agreement of the two methods in terms of absolute values is very low when concentrations are identical (9-24%), increasing to 24-56% when there is a difference of one dilution, to more than 70% for two dilutions and between 92.5-100% for three dilutions. Therefore, when different methods are used, identical MIC values should not be expected; what is necessary is that changes in absolute values do not cause a change in scoring of an isolate as resistant or susceptible, and vice versa, as happened, especially with fluconazole.

In summary, the Alamar colorimetric method is a useful tool in the study of MICs of the five systemic antifungal agents available for systemic treatment of fungal infection. With *C. neoformans* however, the distribution of MICs (fluconazole and 5-flucytosine) does not provide sufficient concordant results in a low but not negligible number of isolates.

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In vitro susceptibility of *Cryptococcus neoformans* serotypes to GM 237354 derivative of the sordarin class

Empfindlichkeit van *Cryptococcus neoformans*-Serotypen *in vitro* für das Sordarin-Derivat CM 237354

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Key words. *Cryptococcus neoformans*, serotypes, antifungal susceptibility, sordarin, amphotericin B. **Schlüsselwörter.** *Cryptococcus neoformans*, Serotypen, Antimycetische, Empfindlichkeit, Sordarin, Amphotericin

SUInInary. *In vitro* susceptibility to the sordarin derivative GM 237354 and amphotericin B were tested in a total of 190 *Cryptococcus neoformans* clinical isolates from different geographical areas of Spain and South American countries. Minimal inhibitory concentrations (MICs) were obtained using the NCCLS reference microbroth dilution method and analysed according the serotypes of *Cr. neoformans*. The MICs for amphotericin B were lower than 1.0 J.1g ml⁻¹ (MIC_{90%} 0.5 J.1g ml⁻¹, MIC_{50%} 0.125 J.1g ml⁻¹) but five isolates showed MICs of 2.0 J.1g ml⁻¹ to GM 237354 (MIC_{90%} 1.0 J.1g ml⁻¹, MIC_{50%} 0.5 J.1g ml⁻¹). *Cryptococcus neoformans* var. *gattii* serotype B, was significantly less susceptible than A and AD serotypes ($P=0.047$ and $p=0.022$, respectively).

ZusalnInenfassung. An 190 klinischen Isola- ten von *Cryptococcus neoformans* aus unterschiedlichen Regionen Spaniens und Südamerikas wurde die Empfindlichkeit *in vitro* für das Sordarin-Derivat GM 237354 im Vergleich zu Amphotericin B getestet. Die MHK-Werte wurden mittels NCCLS- Mikroverdünnungs- Referenzmethode erhoben und ein Vergleich nach *Cr. neoformans*-Serotypen durch- geführt. Die MHKs für Amphotericin B lagen unter

1.0 ~g ml⁻¹ (MHK_{90%} 0.5 ~g ml⁻¹, MHK_{50%} 0.125 ~g ml⁻¹), aber fünf Isolate zeigten MHKs von 2.0 ~g ml⁻¹ für GM 237354 (MHK_{90%} 1.0 ~g ml⁻¹, MHK_{50%} 0.5 ~g ml⁻¹). *Cryptococcus neoformans* var. *gattii* Serotyp. B war signifikant weniger empfindlich als die Serotypen A und AD ($P=0.047$ bzw. $p=0.022$).

Introduction

Sordarin derivatives are a new class of antifungal drugs that appeared to be selective and potent inhibitors of the fungal protein synthesis system [1,2]. Compound GM 237354 have been isolated and characterised from a broth obtained from the culture of the mould *Graphium putredinis*. The broth media inhibited fungal protein synthesis of *Candida albicans* and the following isolation of the active component showed some similarity to sordarin and zofimarin, but the novel antibiotic was designed with the number 237354. In first studies GM 237354 have shown *in vitro* activity against *C. albicans*, *Candida tropicalis*, and *Cryptococcus neoformans* with minimal inhibitory concentrations (MICs) ranging between 0.008 and 1.0 ~g ml⁻¹. In an experimental model of mice with systemic candidosis a therapeutic effect was achieved following subcutaneous injections of 40 mg kg⁻¹ [3].

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As sordarins seem to be promising antifungal drugs, the objective of this study was to analyse a large number of *Cr. neoformans* isolates for their MICs against sordarin derivatives comparing it with amphotericin B and considering the source of yeasts and serotypes of the strains.

Materials and methods

Isolates

A total of 190 clinical isolates of *Cr. neoformans* were included in the study. Clinical samples were mostly recovered from HIV-infected patients with cryptococcosis from different countries including Spain ($n = 117$), Argentina ($n = 29$), Brazil ($n = 22$), and Cuba ($n = 22$). Of these, 175 were from the *neoformans* variety (serotype A 72%, serotype D 11.5%, serotype AD 8.4%) and the remaining 15 belonging to the *gattii* variety, serotype B. Method for serotyping have been previously described [4].

Five reference isolates of *Candida albicans* (A TCC 90028), *Candida krusei* (A TCC 6258), *Candida parapsilosis* (A TCC 220 19) and *Cryptococcus neoformans* (A TCC 90 112 and R VB 20 128) were included for quality control. They were used each time the test was done.

Methodology

For preparing the inoculum the yeast were grown on Sabouraud glucose agar plates for 24 h for *Candida* spp. and 48 h for *Cr. neoformans*. Yeast concentrations were prepared for at least five colonies in sterile saline and adjusted at 0.5 McFarland standard. The inocula were adjusted to a concentration of $1-2.5 \times 10^3$. An aliquot of 0.1 ml was added to each well of the microdilution tray.

Amphotericin B in sterile powder was provided by Squibb Laboratories (Esplugues del Uobregat, Barcelona, Spain); GM 237354 was obtained from GlaxoWellcome, S.A. (Tres Cantos, Madrid, Spain) as sodium salts that were initially solubilized in dimethyl sulfoxide (DMSO) at a concentration of 5 mg ml⁻¹ and then diluted in medium to the appropriate concentration. Solutions were prepared just before use. The microdilution method was carried out and interpreted according to the recommendations of the National Committee for Clinical Laboratory Standards (M27-A document) [5].

RPMI 1640-glutamine plus 2% glucose, in 3-[N-Morpholino] propanesulfonic acid (MOPS) buffer to pH 7 was used as culture medium. Antifungal stock solutions were prepared in water (GM 237354) or DMSO solvent for amphotericin B at concentrations 100 times higher than the working solution. The final concentration was prepared from the antifungal stock solution in RPMI and 2% glucose, with a range between 0.03 and 16 µg ml⁻¹. Antifungal agents were dispensed in sterile, wrapped polystyrene flat-bottom assay plates in wells I to 10 in each row; drug-free medium was dispensed in wells II and 12, the latter serving as a sterility control, and well III as growth control. The

inoculated plates were incubated for 24-72 h at 35 °C. Absorbance were determined with automated plate reader at 414 nm for GM 237354, after agitation of the plates (Mult Labsystem MS, Finland). MIC was defined (lowest concentration in micrograms per millilitre the antifungal that inhibits development of yeast growth in comparison with the growth control (when the antifungal prevents any visible growth). For amphotericin B the MIC was 100% inhibition, when the medium was optically clear. An additional automated test was also carried out at 620 nm.

Analysis of results

The Student's *t*-test for paired data was used for statistical analysis. Statistical significance was $p < 0.05$. The SPSS computer program (SPSS- Chicago, IL, USA) was used for analysis of data.

Results

The MICs of amphotericin B for the control strains agreed with the NCCLS defined values; Repetition of the tests did not give more than a dilution of difference for both antifungals. Ranges of MICs to amphotericin B were equal; lower than 0.03 to 1 µg ml⁻¹, with a geometric mean (SD) of 0.11 (0.17) µg ml⁻¹. The MIC was 0.5 µg ml⁻¹ and MIC_{50%} was 0.125 µg ml⁻¹. For the sordarin derivative GM 237354, the MIC was 0.03 to 2.0 µg ml⁻¹, the highest MICs found in only five isolates, with a mean of (0.37) µg ml⁻¹ and MIC_{90%} and MIC_{500%} were 0.5 µg ml⁻¹, respectively. The distribution of MICs to this product according to the serotype is presented in Table 1.

There were no statistically significant differences between serotypes A and D ($P = 0.9846$), D and AD ($P = 0.3013$), A and AD ($P = 0.2274$) or D and AD ($P = 0.12$). No differences were found according to the geographical origin (Spanish or American) of the isolates. Serotype B of *Cr. neoformans* variety showed significantly higher MIC values than serotype A ($P = 0.0475$) and serotype D ($P = 0.022$).

Discussion

With the use of the standardized method susceptibility testing it is now possible to obtain reproducible and accurate results in order to determine the levels of susceptibility of strains of *Cr. neoformans*. [6, 7].

Table 1. Distribution of MICs ($\mu\text{g ml}^{-1}$) of GM 237354, sordarin antifungal for 190 isolates of *Cryptococcus neoformans* according to serotypes

Serotypes	n	Range	Mean (SD)	MIC _{90%}	MIC _{50%}
A	137	0.003–2.0	0.537 (18.6)	1.0	0.5
B	15	0.5–2.0	0.833 (3.5)	1.0	1.0
D	22	0.06–2.0	0.587 (3.3)	1.0	0.5
AD	16	0.25–1.0	0.5 (4.1)	1.0	0.5

Over recent years, the lack of effective therapy for AIDS patients has prompted research into new antifungal molecules, as in the case of sordarin [2]. In this case the higher activity of this drug against *Pneumocystis carinii* and some dimorphic fungi, increased interest in the product [8, 9].

The *in vitro* activity of GM237354 against *Cr. neoformans* has been investigated in only 20 strains for which the origin and biotype or serotype were not explained [8].

Varieties and serotypes of *Cr. neoformans* are important with regard to ecology and epidemiology and virulence. Genetic analyses resulted in the new variety *grubii* being proposed for the present A serotype [9]. Up to now the correlation of *Cr. neoformans* serotypes with susceptibility or resistance to antifungals has merited little attention [10-13].

In our study, all the isolates showed MIC values for amphotericin B that were lower than 1.0 J.1g ml⁻¹ with MIC_{90%} of 0.5 J.1g ml⁻¹ confirming the virtual absence of resistance of *Cr. neoformans* to this polyene. With GM 237354 in five isolates the MICs presented values of 2 J.1g ml⁻¹, and the MIC_{90%} was 1.0 J.1g ml⁻¹. These values were higher than the results observed in the previous study [8] in which the MIC range was 0.015-1.25 J.1g ml⁻¹ and MIC_{90%} was 0.25 J.1g ml⁻¹. These differences in susceptibility could be explained by the larger number of isolates tested in our study, and by the geographical diversity of strains and serotypes. Moreover, Herreros *et al.* [8] used yeast nitrogen base as the culture medium instead of RPMI 1640.

The clinical significance of the higher MICs observed needs to be determined in an animal model.

When the susceptibility was correlated to serotypes it was found that the MICs associated with the B serotype were higher in comparison with the A and AD serotypes. This tendency should be confirmed with a larger number of *Cr. neoformans* serotype B isolates.

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Cartas al Director

Caracterización molecular de *Cryptococcus neoformans* var. *gattii* causante de brotes epidémicos de criptococosis en cabras

Sr. Director,

Entre 1990 y 1994, en la provincia de Cáceres (España), se produjeron varios brotes de neumonía grave asociada a caquexia que afectó un número variable de cabras -entre el 2,5 y el 12% de las integrantes de los rebaños-. En algunos casos también existía una afectación encefálica y hepática y se registró una elevada tasa de mortalidad de los animales enfermos [1]. La identificación de aislados obtenidos por necropsia de animales, en cinco de los brotes, demostró que el agente etiológico era una levadura capsulada, identificada como *Cryptococcus neoformans*. Posteriormente el análisis bioquímico y serológico de 13 cepas aisladas de seis cabras, que se habían conservado en la colección de la Facultad de Veterinaria de la Universidad de Extremadura, permitió comprobar que correspondían al serotipo B de la variedad *gattii* [2].

La demostración de que *C. neoformans* var. *gattii* existe en nuestro país, y que es capaz de ocasionar infecciones graves en animales sin otras patologías previas, constituye un hallazgo destacable que enriquece la información referida a la epidemiología de la criptococosis.

Para caracterizar las cepas aisladas se procedió a un análisis molecular utilizando la técnica del *Random Amplified Polymorphic DNA* (RAPD). Para ello se efectuó la extracción del ADN de todos los aislamientos mediante la técnica de Lehmann *et al.* [3] modificada. Se efectuó una amplificación por PCR siguiendo la técnica de Williams *et al.* [4] con cuatro diferentes cebadores: CNI (5'TACCCCGCCATA1TCCAT3'), 5S0R (5' ATGGGAA- TACGACGTGCTGTAG 3'), CI (5' ACGGTACTACT 3') y C3 (5'GTITCCGCCCC 3'), estos dos últimos se emplearon de forma simultánea. Para la amplificación se utilizó un termociclador (Perkin-Elmer Cetus, EE.UU.) realizándose 40 ciclos y el material obtenido se analizó por electroforesis en gel de agarosa al 2% con tinción de bromuro de etidio. Después de una detallada observación visual, las imágenes se escanearon para ser procesadas en el sistema computarizado GEL DOC 1000, Molecular analysis (Bio-Rad Laboratories, California USA).

Se determinó el tamaño molecular de los fragmentos de ADN amplificados. La repetitividad de la técnica se confirmó por la amplificación del ADN de un mismo organismo en tres ocasiones diferentes. Los fragmentos de dudosa interpretación no fueron asumidos como elementos de discriminación de los patrones [5,6].

Los datos proporcionados por el análisis de imagen han permitido agrupar las cepas con idéntico perfil electroforético en dos diferentes patrones genotípicos que han sido denominados arbitrariamente A y B. Cuando se disponía de dos aislamientos de un mismo animal, el patrón genómico fue el mismo. El patrón A ha sido el dominante puesto que en todos los brotes se ha demostrado la presencia del mismo; sin embargo, en uno de los episodios ocurrido en la localidad de Vera, en que fue posible disponer de varias cepas aisladas de dos animales diferentes, se encontró el segundo patrón, o B (Tabla I).

Estos resultados sugieren que, con la metodología utilizada, se ha demostrado que en las diferentes zonas de Cáceres donde tuvieron lugar los brotes epidémicos existe un tipo dominante de *C. neoformans* var. *gattii*, sin embargo esta cepa coexiste en la misma área por lo menos con otro tipo genómico.

Los resultados obtenidos, plantean diversas incógnitas como la de determinar el nicho ecológico de esta especie en la naturaleza en Cáceres y en otras partes de la península ibérica y conocer la diversidad genética de las mismas, para poder relacionarla con la capacidad infectiva de esta especie.

T.bla 1 Distribución de KJs aislamientos de *Cryptococcus neoformans* var. *gattii* obtenido de la necropsia de cabras afectadas de criptococosis y el patrón genómico de las mismas determinado utilizando el *Random Amplified Polymorphic DNA*.

Animar	Cepas Muestra	Localidad	Cebador utilizado SSOR			Patrón
			CN1			
				C1+C3		
1	48A	Pulmón Pescueza Pulmón			VI	A
1	49A	Pescueza Pulmón Serradilla			VI	A
2	50A	Pulmón Serradilla Hígado Casas			VI	A
2	60A	de Millan Encéfalo Casas de			VI	A
3	51 A	Millan Pulmón Madroñera			VI	A
3	52A	Pulmón Madroñera Intestino			VI	A
4	58A	Vera			VI	A
4	59A	Pulmón Vera			VI V	A
5	56A	Pulmón Vera			VI	B
6	53A	Pulmón Vera			VI	A
6	54A	Pulmón Vera			VI	A
6	55A				VI	A
6	57A					A

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