

ARTÍCULO ORIGINAL

***Malassezia* yeast species isolated from patients with dermatologic lesions**

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Introduction. Yeasts of the genus *Malassezia* form a normal component of skin flora, but are also associated with several dermatological disorders. Since 1996, the description of new species in this genus have led to new questions about their epidemiology and pathogenicity.

Objective. Herein, the frequency of *Malassezia* species in individuals with pityriasis versicolor, atopic dermatitis, seborrheic dermatitis, seborrheic dermatitis was compared in HIV patients and healthy individuals. Three body sites were selected for examination -head, thorax, and upper and lower extremities.

Material and methods. The 154 *Malassezia* species were isolated from 112 individuals and grouped as follows: 39 with seborrheic dermatitis (20 were HIV-positive patients), 18 with pityriasis versicolor, 18 with atopic dermatitis and 37 without dermatological lesions. HIV patient samples were examined microscopically, and specimens from both patients and healthy subjects were cultured on modified Dixon agar medium. Subsequently, isolates were identified by macroscopic, microscopic and physiological characteristics.

Results. The most commonly isolated species were *Malassezia globosa* (37.5%), *M. sympodialis* (31.3%) and *M. furfur* (31.3%). *Malassezia globosa* predominated in patients with pityriasis versicolor (67%) and in HIV-positive patients with seborrheic dermatitis (85%). In non-HIV patients with atopic dermatitis or seborrheic dermatitis, *M. furfur* and *M. restricta* were isolated in 72% and 26% of the cases, respectively.

Conclusion. Several conclusions were evident. First, *Malassezia* species was present in subjects with and without dermatological pathologies. Second, the species frequency in the sampled population differed from frequencies reported from other geographic areas. Third, *Malassezia globosa* was involved at high frequency in patients with dermatological pathologies, suggesting a higher pathogenicity of this species. Additional studies on each species are recommended to clarify their pathogenic roles in association with HIV-positive and normal subjects.

Keywords: *Malassezia*, tinea versicolor, seborrheic dermatitis, HIV, atopic dermatitis

Especies de *Malassezia* aisladas de pacientes con lesiones dermatológicas

Introducción. Las levaduras del género *Malassezia* forman parte de la flora normal de la piel y se asocian con varios tipos de lesiones dermatológicas. Desde 1996, la descripción de nuevas especies dentro de este género planteó numerosos interrogantes sobre su epidemiología y patogenicidad.

Objetivo. El fin de este trabajo fue encontrar la frecuencia de las diferentes especies de *Malassezia* en individuos con pitiriasis versicolor, dermatitis atópica, dermatitis seborreica, dermatitis seborreica en pacientes positivos para VIH, y en individuos sin lesiones a partir de tres zonas corporales (cabeza, tórax, extremidades inferiores y superiores).

Materiales y métodos. Se identificaron 154 aislamientos de especies de *Malassezia* de 112 individuos: 39 con dermatitis seborreica de los cuales 20 eran positivos para VIH, 18 con pitiriasis versicolor, 18 con dermatitis atópica y 37 individuos sin lesión dermatológica (controles). Las escamas de piel de los pacientes se observaron microscópicamente y las muestras de pacientes y controles se cultivaron en agar Dixon modificado. Posteriormente, se observaron

las colonias y se identificaron según sus características macroscópicas, microscópicas y fisiológicas.

Resultados. Las especies más frecuentemente aisladas en el total de los grupos estudiados fueron: *Malassezia globosa* (37,5%), *M. sympodialis* (31,3%) y *M. furfur* (31,3%). *M. globosa* predominó en los aislamientos de pacientes con pitiriasis versicolor (67%) y en pacientes positivos para VIH con dermatitis seborreica (85%). En pacientes con dermatitis atópica y dermatitis seborreica sin diagnóstico de VIH, se aislaron *M. furfur* y *M. restricta* en el 72% y el 26% de los casos, respectivamente.

Conclusiones. Las diferentes especies de *Malassezia* pueden aislarse de pacientes con lesiones dermatológicas o sin ellas. Además, la frecuencia de especies en la población muestreada difiere de lo reportado en otras zonas geográficas. Se destaca la alta frecuencia de *M. globosa* en pacientes con lesiones lo que podría indicar la mayor patogenicidad de esta especie. No obstante, se requieren estudios adicionales para determinar el papel de las diferentes especies del género *Malassezia*, asociadas con estas patologías.

Palabras clave: *Malassezia*, pitiriasis versicolor, dermatitis seborreica, VIH, dermatitis atópica.

Lipophilic yeasts of the genus *Malassezia* are part of the skin's normal flora which under the influence of certain predisposing factors can give rise to different dermatological disorders (1-3). Nowadays, next to pityriasis versicolor, some authors associate the presence of *Malassezia* yeasts with other dermatological disorders such as atopic dermatitis, seborrheic dermatitis, some forms of folliculitis, psoriasis, Gougerot-Carteaud reticulated confluent papillomatosis, annular centrifugal erythema, seborrheic marginal blepharitis, intertrigo, onychomycosis and neonatal acne. In addition, opportunistic systemic infections of *Malassezia* yeasts have been reported in neonates to whom lipidic infusions have been administered, and also in patients in intensive care units (1-4).

Studies on *Malassezia* cutaneous isolates from healthy humans and animals indicate that *M. sympodialis* is the commonest species in healthy and diseased skin. *M. globosa* is frequently isolated from pityriasis versicolor and seborrheic dermatitis, as well as from healthy skin. *M. restricta*, together with other species, has been associated with dandruff and seborrheic dermatitis of the scalp and face, respectively. Both

M. slooffiae and *M. pachydermatis* are infrequently found on human skin, while *M. obtusa* and *M. furfur* are rarely isolated from these body sites. The species *M. furfur* and *M. pachydermatis* have also been isolated from systemic infections (2,5-7).

Despite reports in 1996 describing four new species for the genus (2), and development of a protocol to identify species by conventional laboratory tests (2,8,9), there have been studies in other countries, not in Colombia, aimed for the determination of species usually found in association to certain pathologies (10-19).

The goal of this study was to determine the relative presence of *Malassezia* species occurring on both patients with dermatologic lesions of atopic dermatitis, seborrheic dermatitis, seborrheic dermatitis in HIV+ patients, pityriasis versicolor, and from individuals with no lesions (healthy controls) on three body sites (head, thorax and lower and upper extremities). As far as we know, this is the first report of *Malassezia* isolated from random patient populations in Bogotá, Colombia.

Materials and methods

Studied population

A total of 112 persons residing in Bogotá, Colombia, were included in the study: 75 with skin lesions, of which 18 corresponded to atopic dermatitis, 39 with seborrheic dermatitis, and 20 of which were HIV-infected patients, 18 had pityriasis versicolor and 37 healthy control individuals. The studied population consisted of 65 men and 47 women, ranging in age from 4 to

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60 years. Individuals were sampled from August 1999 to December 2001. Samples were collected from clinically selected patients who had previously attended dermatological consultation or had been hospitalized in the Simón Bolívar Hospital (Bogotá, Colombia).

Sample-collection and processing

Samples were taken from affected areas by scraping with a disposable scalpel and collecting the scrapings in a sterile Petri dish or in between sterile slides sealed with tape, all of which were transported to the laboratory and processed in less than two hours. A portion of the sample was used for direct examination with 20% KOH plus Parker blue-black ink while the remaining portion was homogenized in 2 ml solution of water plus 0.05% Tween 80 and used for culturing. The contents were vortexed and 0.5 ml were inoculated onto modified Dixon-Agar plates (3.6% malt extract, 0.6% peptone, 2% desiccated ox bile, 1% tween 40 0.2% glycerol, 0.2% oleic acid, 1.2% agar, 0.5 g/L chloramphenicol and 0.5 g/L cycloheximide) (2,8), and then incubated at $32\pm 2^{\circ}\text{C}$ for seven days.

Samples were also taken from control volunteers without lesions and who did not receive oral or topical anti-fungal treatments. No direct examinations were done on these volunteers. Samples were collected according to the body area and the number of sites was proportional to samples taken from patients with dermatologic lesions. Samples were collected by rubbing the area with sterile swabs dampened in 2 ml water plus 0.05% Tween 80, which were introduced in tubes with the same solution. In our first study, we showed that no statistically significant differences occurred between both methods (swabbing and scraping) (20) (García MC, Gutiérrez NE, Sopó L. Abstracts of the 96th General Meeting of the American Society for Microbiology, New Orleans, Louisiana, May 19-23, 1996. Abstract F-31:79). Cultures were processed as described above.

Isolation and identification of colonies compatible with *Malassezia*

Cultures were observed by means of the stereomicroscope and colonies compatible with the genus *Malassezia* were selected. The different

morphotypes (different colony characteristics) observed were subcultured for species identification. The isolates were maintained under liquid nitrogen. The conventional tests described by Guého *et al.*, Guillot *et al.*, and Mayser *et al.*, (2,8,9) were performed in order to confirm the genus and species. The identification scheme used in this study is summarized in figure 1.

Three sampling groups were formed, namely group I corresponding to samples from head (scalp, face and neck), group II were samples from the thorax and group III contained isolates from the extremities (upper and lower).

Results were analyzed by Fisher's exact test.

Results

All samples taken from the 75 patients were positive for *Malassezia* yeasts upon direct examination. Samples taken from the 112 individuals (75 with lesions, 37 healthy controls) yielded 154 isolates, as more than one colony was observed in certain primary cultures. The distribution of the *Malassezia* species isolated is presented in table 1. Six *Malassezia* species could be identified from patients and controls, but 22 isolates could not be identified (*Malassezia* spp.). Three species, *M. globosa* (37.5%), *M. sympodialis* (31.3%), and *M. furfur* (31.3 %) were predominant in the group of isolates as a whole. When separating isolations by groups (patients and controls) a significant difference was noticed for *M. globosa*, which was isolated from 48% of the patients and from 16.2% of the controls ($p=0.003$). Otherwise, the various species were represented in equal amounts in both patients and controls.

The distribution of the *Malassezia* species according to the body site sampled irrespective of the dermatological condition is shown in table 2. In group I, the various species were isolated from both patients and controls. However, with the exception of *M. pachydermatis*, these differences were statistically not significant. On the contrary, when samples in groups II and III were tabulated it was noticed that differences existed according to the species isolated. Thus, in group II, statistically significant differences were seen for *M. globosa* between patients (45.2%) and controls

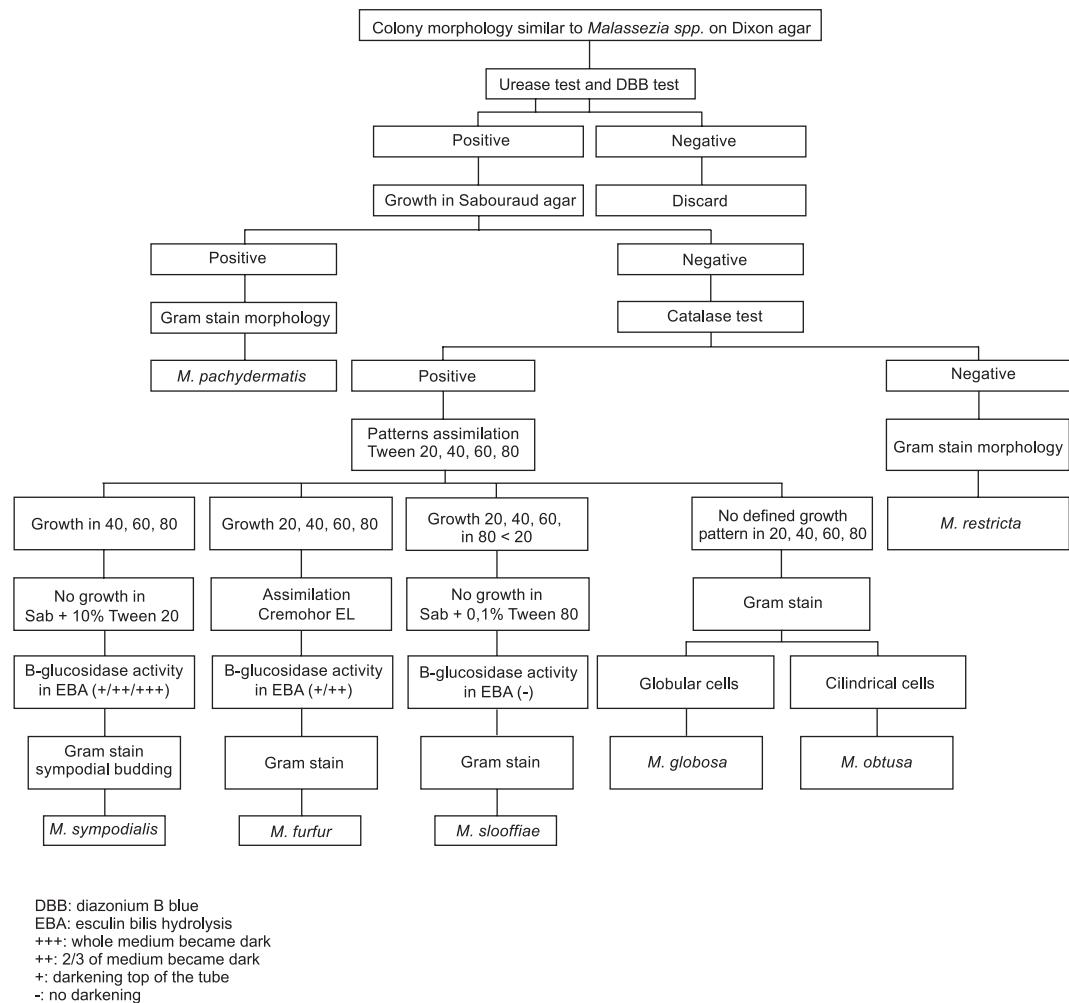


Figura 1. Identification of yeasts compatible with the *Malassezia* genus (modified from references Guého *et al.*, Guillot *et al.*, and Mayser *et al.*

(7.7%) (p=0.035). Inverse figures were noticed for *Malassezia* spp. isolates, namely 38.5% in controls vs. 9.7% in patients (p=0.035). In group III no statistically significant differences occurred for *M. pachydermatis* and *Malassezia* spp. isolated from both groups patients and controls. The *Malassezia furfur* isolates were obtained only from patients, but the small number of samples in this group (n=14) precludes proper interpretation. It can be seen that the largest number of species were isolated from group I, namely 6 identified species and non-identified *Malassezia* spp.

We attempted to determine the role of the *Malassezia* species in the pathogenesis of the

Table 1. Distribution of *Malassezia* species from patients and controls.

| Species | Patients n=75* | | Controls n=37* | | Total n=112 | |
|-------------------------|-------------------|------|-------------------|------|----------------|------|
| | n | % | n | % | n | % |
| <i>M. globosa</i> | 36 | 48,0 | 6 | 16,2 | 42 | 37,5 |
| <i>M. sympodialis</i> | 24 | 32,0 | 11 | 29,7 | 35 | 31,3 |
| <i>M. furfur</i> | 27 | 36,0 | 8 | 21,6 | 35 | 31,3 |
| <i>Malassezia</i> spp. | 12 | 16,0 | 10 | 27,0 | 22 | 19,6 |
| <i>M. restricta</i> | 11 | 14,7 | 2 | 5,4 | 13 | 11,6 |
| <i>M. obtusa</i> | 4 | 5,3 | 1 | 2,7 | 5 | 4,5 |
| <i>M. pachydermatis</i> | 0 | 0,0 | 2 | 5,4 | 2 | 1,8 |

*In some cases, more that one species was isolated from the same sample.

Table 2. Distribution of *Malassezia* species isolated from head, thorax, lower and upper extremities of patients and controls.

| Species | Group I n*=97 | | | | Group II n*=44 | | | | Group III n*=14 | | | |
|-------------------------|---------------|------|----------|----|----------------|------|----------|------|-----------------|------|----------|------|
| | Patients | | Controls | | Patients | | Controls | | Patients | | Controls | |
| | n | % | n | % | n | % | n | % | n | % | n | % |
| <i>M. sympodialis</i> | 15 | 20,8 | 4 | 16 | 7 | 22,6 | 6 | 46,2 | 2 | 18,2 | 1 | 33,3 |
| <i>M. furfur</i> | 14 | 19,4 | 7 | 28 | 6 | 19,4 | 1 | 7,7 | 7 | 63,6 | 0 | 0 |
| <i>M. globosa</i> | 21 | 29,2 | 5 | 20 | 14 | 45,2 | 1 | 7,7 | 1 | 9,1 | 1 | 33,3 |
| <i>Malassezia</i> spp. | 8 | 11,1 | 5 | 20 | 3 | 9,7 | 5 | 38,5 | 1 | 9,1 | 0 | 0 |
| <i>M. restricta</i> | 10 | 13,9 | 2 | 8 | 1 | 3,2 | 0 | 0 | 0 | 0 | 0 | 0 |
| <i>M. obtusa</i> | 4 | 5,6 | 1 | 4 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| <i>M. pachydermatis</i> | 0 | 0 | 1 | 4 | 0 | 0 | 0 | 0 | 0 | 0 | 1 | 33,3 |
| Total of isolations | 72 | | 25 | | 31 | | 13 | | 11 | | 3 | |

Group I: head samples; group II: thorax samples; group III: lower and upper extremities samples

n*: number of isolates of *Malassezia* spp. from each group (I,II,III)

n: number of isolates of *Malassezia* spp. from three body sites of patients and controls

various skin disorders. The species isolated from patients and controls were compared according to the different dermatologic entities. In atopic dermatitis, statistically significant differences were seen for *M. furfur* between patients (13/18) (72%) and controls (8/37) (22%) ($p=0.0002$). Additionally, significant differences were observed in seborrhoeic dermatitis for *M. restricta* in controls (2/37) (5.4%) and patients (5/19) (26%) ($p=0.016$), in HIV-positive patients *M. globosa* was isolated from 17/20 (85%) of the patients vs. 6/37 (16%) of the controls ($p<0.01$), and *M. restricta* occurred in 5/20 (25%) patients and 2/37 (5.4%) controls ($p=0.022$). In pityriasis versicolor significant differences were noticed for *M. globosa* between patients (12/18) (67%), and controls (6/37) (16%) ($p<0.01$).

Discussion

The role of these lipophilic *Malassezia* yeasts in skin problems is controversial, although a number of studies have been undertaken to investigate the ecology and the epidemiology of the different *Malassezia* species in these pathological disorders, as well as of their presence on healthy skin (1-4, 10-19). New questions have been raised about the significance and relationships of the *Malassezia* species in clinical infections (1,4), this is due to factors such as the recent description of new species (2,21,22), the isolation of *Malassezia* as part of the microbial community of human skin rich in sebaceous glands (1,2,6,16), and the

variability in colonization according to different age groups and sebaceous glands activity (15,19,23).

Our results indicate that the most frequently isolated species from all populations studied were *M. globosa* (37.5%), *M. sympodialis* (31.3%) and *M. furfur* (31.3%). In accordance with previous reports, the remaining species *M. restricta*, *M. obtusa*, and *M. pachydermatis* were isolated less frequently (10,11,15,18).

We found isolates that did not correspond to previously reported patterns in the standard identification protocols (2,8,9), and these strains were grouped as *Malassezia* spp., a category that has been reported in previous studies as well (10,23,24). Recently, other new *Malassezia* species have been reported, namely *M. japonica* (21) and *M. dermatis* (22), and these may be similar to some of the unclassified species referred above.

Our results reveal the predominance of *M. globosa* (67%) in patients with pityriasis versicolor ($p<0.01$), which is in accordance with reports from various authors (10-12,17). Many reports consider that *M. globosa* is the etiologic agent of pityriasis versicolor isolated from 25-97% of the cases (6, 9-12,17,18). However, others have reported *M. sympodialis* as the most frequently isolated species (40-62.7%) (14,15).

Moreover, we have found different implications of *Malassezia*-related diseases in HIV-infected

patients, in which the main pathology is not pityriasis versicolor, but seborrhoeic dermatitis with *M. globosa* as the predominant etiologic agent (85%) ($p < 0.01$), followed by *M. restricta* (25%) ($p = 0.022$). In contrast, in a previous study conducted in our laboratory, *Malassezia* spp. colonization was evaluated in HIV-infected patients from the same hospital, *M. restricta* (46%) was found to be the prevailing species (13). The difference could be explained by the sampling period and year, and because of the cutaneous microbial communities occurring on these patients (15,19). Nevertheless, the results of this study are interesting and indicate a superior pathogenic capacity for *M. globosa*, which call for additional study in this group of HIV-infected patients.

M. furfur isolated in 72% of the patients with atopic dermatitis has statistically significant differences ($p = 0.002$), a finding that is also of interest. Although few studies have been conducted in this respect, our data are similar to those of Nakabayashi *et al.* in Japan (10) and differs from others, which report *M. sympodialis* as the most frequently isolated species (53%) in Canada (15) and 40% in Sweden (Faergemann J, Sandstrom M, Back O, Scheynius A, Samhult T, Linder M. Abstracts of the 15th Congress of the International Society for Human and Animal Mycology, ISHAM. San Antonio, Texas, May 25-29, 2003, Abstract 368:459). However, the heterogeneous nature of this disease has been recognized and it has been suggested that *Malassezia* yeasts act as allergens than as etiological agents (1,6,25).

Significant differences were found ($p = 0.016$) in the isolation of *M. restricta* (26%) from patients with seborrhoeic dermatitis which is in accordance with studies performed in Spain where *M. restricta* was isolated in 43%-63.9% of the patients evaluated (11,12). Once more, this differs from other reports, as follows: *M. globosa*, 45% in Canada (15); *M. sympodialis*, 38.5%, and *M. slooffiae*, 34.6% in Mexico (18); *M. furfur* and *M. globosa* in equal percentages (21%) in Japan (10); *M. obtusa*, 35%, and *M. sympodialis*, 30% in other group of patients in Sweden (Faergemann J, Sandstrom M, Back O, Scheynius A, Samhult T, Linder M. Abstracts of the 15th Congress of the International Society for Human and Animal Mycology, ISHAM. San

Antonio, Texas, May 25-29, 2003, Abstract 368:459).

We also evaluated (table 2) the relation between the isolated species and the body sites head, thorax and upper and lower extremities, groups (I, II and III). Our results differ from other reports in the distribution of the species on body sites in different geographic areas of Spain, Japan and Canada (10-12,15,19). These differences are probably related to the ethnic origin of the population, the corporal zones, and the age of the individuals tested, as suggested by several authors (4,15,19). Our findings suggest the need of additional studies to determine the ecology of the different species in healthy skin and the clinical significance of the dermatoses associated with these yeasts, as well as the urgency of new studies to evaluate the differences in susceptibility to antifungal compounds and intra-specific differences using molecular techniques.

Conflict of interests

The authors declare that they have no competing interests.

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