

Bacillus cereus strain isolated from *Demodex folliculorum* in patients with topical steroid-induced rosaceiform facial dermatitis*

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DOI: <http://dx.doi.org/10.1590/abd1806-4841.20165214>

Abstract: The aim of the study was to identify *Bacillus* species from the *Demodex folliculorum* of patients with topical steroid-induced facial rosaceiform dermatitis. Of the 75 patients examined, 20% had clinical spinulosis, while 18.66% had dermoscopic features of *Demodex*: follicular plugs and tails. Of the 17.33% positive patients identified upon microscopy for *Demodex*, samples for bacterial culture were plated on trypticase soy Columbia agar. Identification was performed by microorganisms grown method mass spectrometry. We identified a strain of *Bacillus cereus*.

Keywords: Dermoscopy; Facial dermatoses; Mass spectrometry; Mite infestations; Receptors, steroid; Rosacea

INTRODUCTION

Bacillus oleronius isolated from *Demodex folliculorum* has been identified as a trigger of inflammation in rosacea.¹ Skin lesions associated with an abnormal increase in *Demodex* mites, classified as secondary demodicosis, occur mostly in patients undergoing treatment with topical steroids or calcineurin inhibitors.²

The aim of the study was to assess the microbioma in patients with topical steroid-induced facial rosaceiform dermatitis (TSIFRD) and the possible association between *Bacillus* and *Demodex* species.

METHODS

Patients with TSIFRD were included in this study, entailing clinical examination, dermoscopy and microscopy analysis of *Demodex* species. Furthermore, skin biopsies were performed on patients with inflammatory lesions. For the microscopic examination, samples were harvested from the affected area using a piece of Scotch transparent tape measuring 2cm, which was subsequently displayed in a microscope slide and examined under optical microscopy (10x- 40x Zeiss). The sample for bacterial culture was spun two minutes and plated (trypticase soy agar and Columbia agar with 5% sheep's blood (Oxoid – Thermo Fischer, UK), then incubated at 37°C for 48 hours. Identification was performed by micro-

organisms grown method MALDI-TOF mass spectrometry (Matrix Assisted Laser Desorption Time-of-Flight ionization - Bruker Daltonik GmbH, Bremen, Germany). Colonies were transferred, emulsified and centrifuged, the supernatant was removed and the residue reconstituted in 70% formic acid. Measurements were performed using MALDI-TOF MS Microflex LT through the FlexControl software. The spectra were collected and analyzed via the Biotyper database (database [DB] -5627 Species list). Scores of above 2 were considered significant for genus and species identification. Mass spectrometry was used to identify the genus and species. The local ethics committee approved the study.

RESULTS

Seventy-five patients with variable TSIFRD subtypes were examined. Dermoscopy revealed: telangiectasias (100% of patients), pustules (80%), scales (54.66%) and atrophy (20%), as white structureless patches between vessels. Clinical spinulosis was found in 15 patients (20%), of which 14-18.66% bore dermoscopic features of *Demodex*: follicular plugs and *Demodex* tails (Figures 1 and 2). A biopsy performed on one spinulosis area showed a histopathologically characteristic image (Figure 3). Thirteen patients (17.33%) were positive for *Demodex* upon microscopy: 8 for *D. folliculorum* and 5

Received on 01.10.2015

Approved by the Advisory Board and accepted for publication on 11.02.2016

* Work performed at the University Dunarea de Jos, Faculty of Medicine and Pharmacy; Central Reference Laboratory Synevo, University of Medicine and Pharmacy Carol Davila – Bucharest, Romania.

Financial support: Acknowledgements: This paper is supported by the Sectoral Operational Programme Human Resources Development (SOP HRD), financed by the European Social Fund and by the Romanian Government under the contract POSDRU/159/1.5/S/137390/

Conflict of interest: none

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FIGURE 1: Spinulosis of the face



FIGURE 2: Dermoscopic follicular plugs and Demodex tails

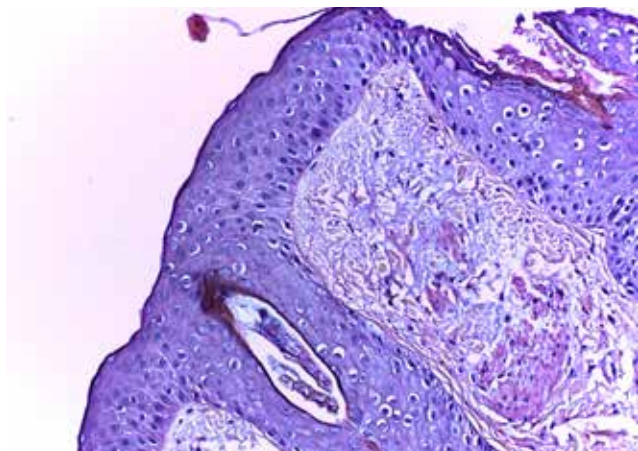


FIGURE 3: Histopathology of spinulosis revealed Demodex folliculorum; hematoxylin/eosine staining,40x

for *D. brevis* (Figures 4 and 5). Among the 8 patients with *D. folliculorum*, one had positive *Bacillus cereus* cultures. No *B. oleronius* or any other bacillus species were found. An analysis of the macroscopic diagnosis of bacterial cultures of the *Bacillus* species oriented morphology *B. cereus* colonies, which were dull gray and opaque, with a rough matted surface. The genus and species were identified by mass spectrometry (Figures 6 and 7).

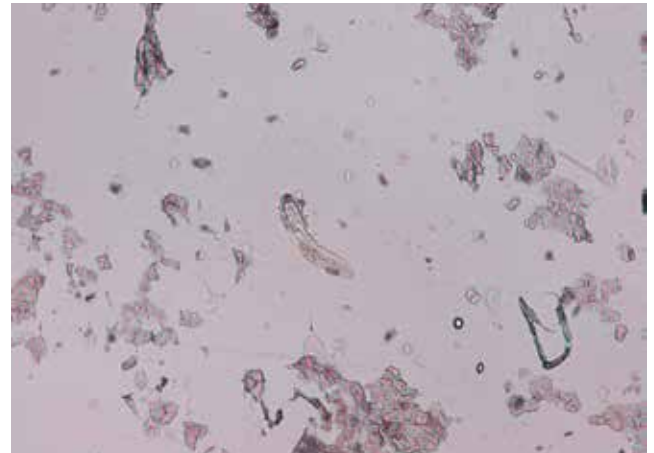


FIGURE 4: Demodex folliculorum smear,10x



FIGURE 5: Demodex brevis smear,20x

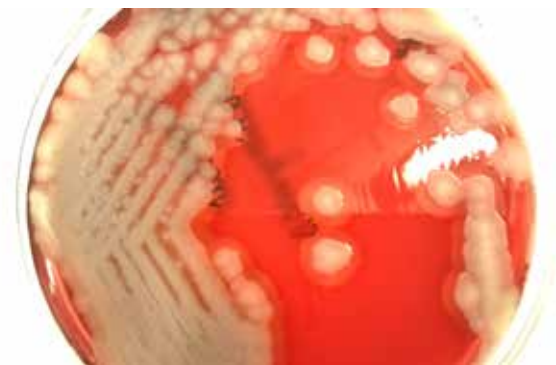


FIGURE 6: Bacillus cereus culture at 72 hours on Colombia agar with 5% sheep's blood

DISCUSSION

B. oleronius was isolated from only one dissected *Demodex* harvested from one patient with papulopustular rosacea.³ The absence of serum reactivity to *Bacillus* spp. antigens in 20% of patients with initial rosacea, along with the presence of antibodies in 40% of the controls without visible rosacea (as recorded in previous studies), raises questions about the role of these bacteria as atrig-

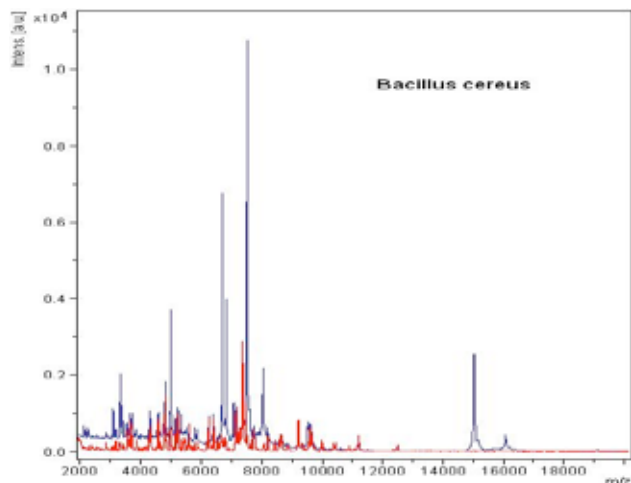


FIGURE 7: The spectrum obtained from the identification of *Bacillus cereus* by MALDI-TOF spectrometry

ger of the post-steroid rosacea inflammatory process.⁴ *B. cereus* is a Gram-positive, aerobic-to-facultative, spore-forming rod widely distributed environmentally, bearing close phenotypic and genetic (16S rRNA) links to several other *Bacillus* species, especially *B. anthracis*.⁵ One reservoir for *B. cereus* is the intestinal tract of invertebrates. It has been observed in the guts of certain arthropods which is regarded as the normal intestinal stage in soil-dwelling insects attached to the arthropod intestinal epithelium, where they sporulate.^{6,7} In the food industry, *B. cereus* spores are particularly

troublesome because spores can be refractory to pasteurization and gamma radiation, while their hydrophobic nature allows them to adhere to surfaces.⁸ *B. cereus* spores may colonize skin on the hands and feet, which are often in contact with the environment through microscopic skin abrasions. The pathogenicity of *B. cereus* is intimately associated with tissue-destructive or reactive exoenzyme production: four hemolysins, three phospholipases, one emesis-inducing toxin, and three enterotoxins: hemolysin BL (HBL), nonhemolytic enterotoxin (NHE) and cytotoxin K.⁹ In 2008, Dohmae *et al.* described a *B. cereus* nosocomial infection from reused towels in Japan.¹⁰ This study focused on the correlation between *D. folliculorum* and types of *Bacillus* species. The low density of *Demodex* species in the patients included in the study could explain the lack of identification of *B. oleronius* culture, which may underline the difference of microbiome between typical rosacea and TSIFRD.

CONCLUSIONS

In a series of 75 patients with topical steroid-induced facial rosaceiform dermatitis (TSIFRD), 15 had clinical spinulosis, 14 had dermoscopic features of *D. folliculorum* - follicular plugs, *Demodex* tails confirmed by scraping, microscopy and histopathology. Eight out of 15 patients had *D. folliculorum*, and one case revealed associations with positive *B. cereus* cultures; while no *B. oleronius* was isolated. These results show a possible difference between the microbiome found in TSIFRD and previously reported data on microbiome in rosacea. Larger studies are needed to confirm correlations between *Demodex* and *Bacillus* species in different forms of rosacea and their implication in the inflammatory process. □

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How to cite this article: Tatu AL, Ionescu MA, Clatici VG, Cristea VC. *Bacillus cereus* strain isolated from *Demodex folliculorum* of patients with topical steroid induced rosaceiform facial dermatitis. *An Bras Dermatol.* 2016;91(5):676-8.