PS2: 44

Identification of black aspergilli group based on a polyphasic approach, including MALDI-TOF ICMS

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Black aspergilli is a group of filamentous fungi comprised by species of Aspergillus section Nigri. They can be isolated from different environments however the main habitat of these species is the soil. According to Samson et al. (2007) there are 19 species of Asperaillus section Niari accepted. The species identification must be delineated based on a polyphasic approach, including morphology, physiology, profile of secondary metabolites and molecular biology (Samson and Varga, 2009). Additionally, according to Santos et al. (2010a, 2010b) it is clear that spectral analyses add value to the polyphasic approach. Matrix-Assisted Laser Desorption Time-Of-Flight Intact Cell Mass Spectrometry (MALDI-TOF ICMS) is a spectral technique that analyses the chemical molecular mass of the microbial cellular composition providing rapid and discriminatory fingerprints for identification. This work aimed to perform a polyphasic approach based on morphological, biochemical and spectral analysis by MALDI-TOF ICMS for the characterisation and identification of the section Nigri. Seventy-four isolates of the section Nigri deposited at University of Recife Mycology (URM) Culture Collection were analysed. Additionally, 12 type strains of the section Nigri deposited at Micoteca da Universidade do Minho (MUM) Culture Collection were used as reference for MALDI-TOF ICMS studies. The data obtained from the polyphasic approach indicates that MALDI-TOF ICMS results corroborate with those data obtained using classical taxonomy and biochemical analyses. Overall. from the 74 cultures, 75% were finally identified as A. niger, 15% as A. japonicus, 5% as A. carbonarius, 4% as A. aculeatus and 1% as A. foetidus. Moreover, the biochemical analyses showed that from the whole population of A. niger 20% and 13% were characterised as ochratoxin A (OTA) and fumonisin B₂ producers, respectively. A. carbonarius and A. foetidus were in total OTA producers. The 74 isolates belonging to section Niari deposited at URM were deeply studied and their associated information update and regualified.

Samson et al (2007). Stud. Mycol. 59: 129-145. Samson et al (2009). Med. Mycol. 1-8.Santos et al (2010a). J. App. Microbiol. 108: 375-385. Santos et al (2010b). Res. Microbiol. 161: 168-175. 202