TITLE: MOLECULAR IDENTIFICATION OF PARACOCCIDIOIDOMYCOSIS AGENTS

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ABSTRACT:

Paracoccidioidomycosis (PCM) is a systemic fungal infection caused by etiologic agents belonging to the Paracoccidioides genus. The disease is acquired following inhalation of Paracoccidioides propagules present in the soil, leading to infections varying between acute to chronic PCM. The disease has a wide distribution in Latin America, being Brazil responsible for ~80% of the cases. Laboratory diagnosis of PCM is traditionally made by combining direct exam, culture, and serological assays. However, these techniques are laborious and time-consuming. To overcome this problem, we propose a PCR-based assay to, not only identify Paracoccidioides, but also differentiate between the biological species P. brasiliensis complex and P. lutzii, thus far contributing to achieving rapid and accurate diagnosis of PCM. The 43,000-Da glycoprotein (gp43/ exon 2) sequences from P. brasiliensis and P. lutzii were retrieved from GenBank and used for in silico screening to identify parsimony informative regions (that were divergent interspecifically and conserved intra-specifically), which could be used for primer design. Two species-specific pairs of primers were designed, one for each Paracoccidioides. Primers candidates were validated in silico using Primer-BLAST software and no amplicon was obtained from other pathogenic or non-pathogenic fungi (Fungi taxid4751), but Paracoccidioides. These pairs of primers were tested in vitro as singleplex PCRs using 48 P. brasiliensis and 13 P. lutzii isolates, with high specificity (100%) and sensitivity (100%). Judging from the different amplicon sizes (308 bp for P. brasiliensis and 142 bp for P. lutzii), we optimized a duplex PCR-assay combining both pairs of primers, to identify Paracoccidioides agents in a single round reaction. All samples evaluated were positive for each species-specific pairs of primers using a duplex-PCR assay, which matched the results obtained by the DNA barcoding ITS1/2+5.8s region amplification and sequencing (very good agreement; Kappa value = 1.0). Our duplex-PCR assay showed the applicability of this method, which exhibited an absolute specificity (100%) for the two species and a high sensitivity (100%) even down to low DNA concentrations (1 pg) in a single round reaction. We demonstrated that our PCR-assay has the potential to aid the diagnostic of PCM agents with high accuracy and efficiency, being an inexpensive and fast molecular tool.

Keywords: *Paracoccidioides brasiliensis, Paracoccidioides lutzii,* paracoccidioidomycosis, molecular diagnosis, duplex PCR-assay

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