

A fast and efficient method for total DNA extraction from soil filamentous fungi

*Un método rápido y eficiente para la extracción de ADN
total de hongos filamentosos del suelo*

Wilson Huanca-Mamani^{1*}, Ricardo Salvatierra Martínez¹,
Germán Sepúlveda-Chavera¹

ABSTRACT

Extraction of high quality of DNA from fungus can be affected because the presence of the complex cell wall, high content of polysaccharides or secondary metabolites. In this research, we adapt a simple method of DNA extraction and purification of from fungus isolated from soil for Polymerase Chain Reaction (PCR) applications. The methodology described is both rapid and cost effective.

Key words: DNA extraction, fungi, soil, PCR.

RESUMEN

La extracción de ADN de alta calidad de hongos puede ser afectada debido a la presencia de una pared celular compleja, alto contenido de polisacáridos o metabolitos secundarios. En esta investigación se adaptó un simple método de extracción y purificación de ADN de hongos filamentosos aislados del suelo, para aplicaciones tipo Polymerase Chain Reaction (PCR). La metodología descrita es rápida y de bajo costo.

Palabras clave: extracción de ADN, hongos, suelo, PCR.

Introduction

Some soil filamentous fungi, like *Fusarium*, among other infecting a wide range of crop plants giving rise significant quantitative and qualitative losses to global agriculture (Mishra *et al.*, 2003). The accurate identification of these fungi is difficult because has traditionally been based according to morphological characters (Chen *et al.*, 2011).

The molecular methods has emerged as a possible answer to problems associated with the fungal genetics diagnostics. Polymerase chain reaction (PCR) is a technique widely used in fungal research (Cenis, 1992) mainly for the identification of fungi, since it gives more validity to fungi identification based on both, phenotypic and genotypic attributes (Guarro *et al.*, 1999; Ge *et al.*, 2011). The most relevant methods used in the molecular identification of fungi are based on the

variability of the ribosomal genes (White *et al.* 1990; Hibbett, 1992), as the internal transcribed spacer (ITS) (Gardes and Bruns, 1993). However, fungi possess cell walls made of complex structures such as chitin, glucans lipids and other polymers, which rupture of the cell wall and draw-quality DNA, which is one of the biggest challenges to generate acceptable PCR results (Karacousisa *et al.*, 2006). There are several DNA extraction methods and the most of then are complex protocols and can may require up to several hours. Recently, Chen *et al.*, (2011) reported a rapid method for fungal DNA extraction method, adapted from Chomczynski and Rymaszewski (2006), developed to bacterial DNA samples.

In this research, we adapt a rapid and efficient DNA extraction method, initially described in bacteria (Atashpaz *et al.*, 2010) for species of soil filamentous fungi.

¹ Facultad de Cs. Agronómicas, U. Tarapacá. Avda. General Velásquez 1775, Arica, Chile.

* Autor para correspondencia: whuanca@uta.cl

Materials and Methods

Fungal material

The method was tested in the extraction of DNA of 4 species of soil filamentous: *Trichoderma sp.*, *Fusarium oxysporum*, *Penicillium sp.*, *Paecilomyces sp.* donated by the Plant Pathology group, Faculty Agronomic Sciences of the Tarapacá University, Chile, which were obtained from different soil in the Azapa Valley in Northern Chile (18° 28'60 "S, 70° 13'60" W). Pure cultures of each species were maintained on potato dextrose agar (PDA) at 4° C until use.

DNA extraction procedure

Selected fungal were grown PDA. Three small lump of mycelia is added into a 1.5 mL Eppendorf tube by using a sterile toothpick and ground with pestle to disrupt the micelia. Then, 500 µL lysis buffer (2% CTAB, 100 mM Tris-HCl, 1.4 M NaCl, 1% PVP, 20 mM Na₂EDTA, 0.2% LiCl. The pH was adjusted at ~8) was added and ground with pestle again and incubate at 65 °C per at least 30 min. The mixture was centrifuged at 6,160 g (10,000 rpm) for 5 min at room temperature (RT). The supernatant was transferred into a new tube and an equal volume of chloroform-isoamylalcohol (24: 1 vol/vol) was added. Then the tube was gently flipped several times and centrifuged at 8,870.4 g (12,000 rpm) for 10 min at RT. The upper phase was then transferred into a new tube. An equivalent volume of cold (-20 °C) isopropanol was added drop-wise to precipitate the DNA (optional, store the mixture at -20 °C for 30 min). The sample was precipitated at 12,073.6 g (14,000 rpm) for 10 min at RT. The supernatant was removed and 500 µL of 70% ethanol (4 °C) was added to the sample and centrifuged at 8,870.4 g (12,000 rpm) for 5 min. The supernatant was discarded and the pellet was dried at (RT). The pellet was dissolved in 50 µL H₂O and DNA solution was stored at -20 °C.

PCR condition

Each total DNA was amplified by PCR in a final volume of 20 µL. The PCR reactions were performed using the ITS primers reported by White *et al.* (1990) (IST3; 5'-GCATCGATGAAGAACGCAGC-3' and ITS4; 5'-TCCTCCGCTTATTGATATGC-3'). Each

reaction contained 2 µL of fungal DNA extraction, 10 pmoles of each primer, 2.5 mM of each dNTP, 2 mM MgCl₂, 1X PCR buffer (KCl), 1 unit of Taq DNA polymerase (Thermo Scientific) and sterile distilled water. Cycling conditions were: 5 min at 94 °C; 35 cycles of 30 sec at 94 °C; 30 sec at 55 °C; 30 sec at 72 °C and final elongation step of 2 min at 72 °C. PCR blank reactions controls were incorporated. Three µL of each PCR product was visualised on 1.5% agarose gels stained with gel-red (Biotium).

Results and Discussion

Most common problems identified for the isolation and purification of DNA from fungi isolated from soil include degradation of DNA owing to endonucleases, high production of polysaccharides and the presence of inhibitor compounds such as secondary metabolites that can reduce yield and purity by binding covalently with the extracted DNA (Möller *et al.*, 1992).

In this research we adapted a rapid and efficient DNA bacterial method for fungi isolated from soil (Atashpaz *et al.*, 2010). This short protocol combines inactivation of proteins with precipitation of acidic polysaccharides by hot CTAB in presence of high salts and only a single selective precipitation of DNA with isopropanol and it is appropriate for simultaneous processing of many samples because all steps can be done in Eppendorf tubes. The growth of mycelia on Petri dishes eliminates the need for still of shaking liquid cultures. The whole procedure can be completed within an hour and is applicable to wide variety of soil fungi including *Trichoderma sp.*, *Fusarium oxysporum*, *Penicillium sp.* and *Paecilomyces sp.* Figure 1 shows the results of the separation of the extracted DNA on a 1% agarose gel, stained with Gel-Red (Biotium) and visualized under UV light. The methods here reported yields high-quality DNA, which is transparent, non-viscous, lacks visible contamination of RNA and non-degradation DNA was observed.

DNA yields obtained using this method ranged 400 to 900 ng/µL with an A₂₆₀/A₂₈₀ ratio close to 1.9 and A₂₆₀/A₂₃₀ around 1.7 indicating very little contamination of proteins, polysaccharides or aromatic compounds and may be used for further PCR analysis.

In order to check the efficiency and reliability of this method, the 18S rRNA genes of the ITS

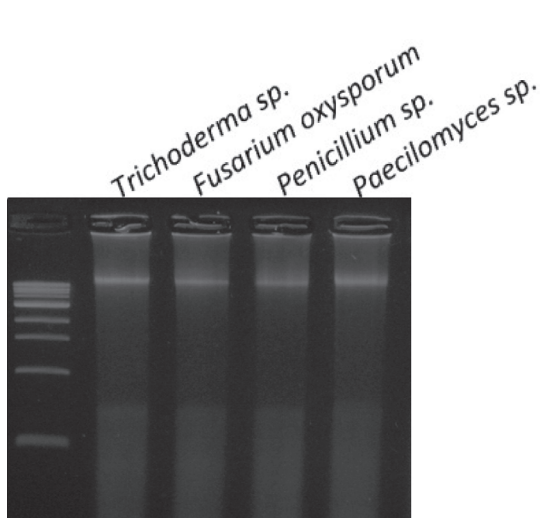


Figure 1. Total genomic DNA extracted from several fungi isolated from soil. DNAs were visualised in 1% agarose gel and stained with Gel-Red. M: 1 Kb DNA Ladder.

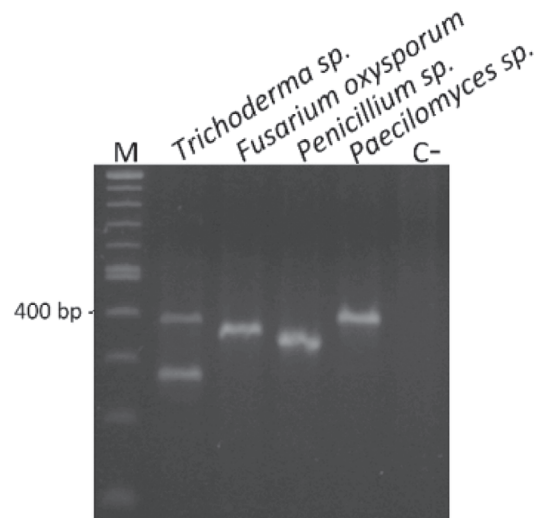


Figure 2. DNA amplification by PCR of the ITS products obtained with ITS3/ITS4 primers. PCRs products from several fungi isolated from soil were visualised in 1% agarose gel and stained with Gel-Red. M: 100 pb DNA Ladder. C-: Negative control.

region were amplified by PCR (Figure 2). Using the ITS3/ITS4 primers, a single band of approximately between 300 and 400 bp is obtained for all strain, however an additional band of approximately 250 bp is observed in *Trichoderma sp.* (Figure 2). It is likely that this robust procedure could be applied to the examination of many other fungal cultures and, possibly, clinical specimens. It provides a rapid, reliable, and low cost alternative to the existing

DNA purification protocols used in molecular biology research.

Acknowledgments

This research was supported by Convenio de Desempeño UTA-MECESUP-2, Foundation for Agrarian Innovation (FIA) PYT-Project 2012-0024 and Mayor-UTA Project Mayor-9711-13.

Literature Cited

- Atashpaz, S.; Khani S.; Barzegari, A.; Barar, J.; Vahed, S.Z.; Azarbaijani, R.; Omid, Y.
2010. A robust universal method for extraction of genomic DNA from bacterial species. *Mikrobiologija*, 79: 562-566.
- Cenis, J.L.
1992. Rapid extraction of fungal DNA for PCR amplification. *Nucleic Acids Research*, 20: 2380.
- Chen, Y.; Prior, B.A.; Shi, G.; Wang, Sh.
2011. A rapid PCR-Based approach for molecular identification of filamentous fungi. *J. of Microbiol.*, 49: 675-679.
- Gardes, M.; Bruns, T.D.
1993. ITS primers with enhanced specificity for basidiomycetes-application to the identification of mycorrhizae and rusts. *Mol. Ecol.*, 2: 113-118.
- Ge, Y.P.; Wang, L.; Lu, G.X.; Shen, Y.N.; Liu, W.D.
2012. A simple and reliable PCR-restriction fragment length polymorphism assay to identify *Candida albicans* and its closely related *Candida dubliniensis*. *Brazilian Journal of Microbiol.*, 43: 873-879.
- Guarro, J.; Gene, J.; Stchigel, A.M.
1999. Developments in fungal taxonomy. *Clin. Microbiol. Rev.*, 12: 454-500.
- Hibbett, D.S.
1992. Ribosomal RNA and fungal systematics. *Trans. Mycol. S C. Japan*, 33: 533-556.
- Karakousisa, A.; Tana, L.; Ellish, D.; Alexioub, H.; Wormald, P.J.
2006. An assessment of the efficiency of fungal DNA extraction methods for maximizing the detection of medically important fungi using PCR. *J. Microbiology Methods*, 65: 38-48.
- Mishra, P.K.; Fox, R.T.; Culham, A.
2003. Development of a PCR-based assay for rapid and reliable identification of pathogenic *Fusaria*. *FEMS Microbiol Lett.*, 218: 329-332.
- O'Donnell, K.; Cigelnik, E.; Casper, H.H.
1998. Molecular phylogenetic, morphological and mycotoxin data support re-identification of the quorn mycoprotein fungus as *Fusarium venenatum*. *Fungal Genet. Biol.*, 23: 57-67.

White, T.J.; Bruns, T.D.; Lee, S.; Taylor, J.W.

1990. Amplification and direct sequencing of fungal ribosomal RNA genes for phylogenetics. *PCR Prot Cols: A Guide to Methods and Applications*, pp. 316-317. *In*: M.A. Innis, D.H. Gelfand, J.J. Sninsky, and T.J. White (eds). Academic Press, Inc., Harcourt Brace Jovanovich, New York, NY, USA. pp. 461.

Möller, E.M.; Bahnweg, G.; Sandermann, G.; Geiger, H.H.

1992. A simple and efficient prot Col for isolation of high molecular weight DNA from filamentous fungi, fruit bodies, and infected plant tissues. *Nucleic Acids Research*, 20: 6115-6116.