

RNA extraction from mycotoxigenic filamentous fungi

Applied Mycology Group, Cranfield Health, Cranfield University, Cranfield, Bedfordshire, UK



CONTEXT

Filamentous fungi cell wall and endogenous RNase activity often hindered the study of the **molecular mechanisms behind secondary metabolites**. **Toxicogenic secondary metabolites** affect human health and represent a risk in the food chain system. The conditions in which **mycotoxins** are produced are still not completely understood especially the molecular mechanism that trigger their production.

The extraction of **high quality RNA in good amounts** was therefore critical to further evaluate the conditions in which mycotoxigenic gene clusters are activated. **Aspergillus flavus** NRRL 3357 was used to evaluate the RNA extraction protocol. A manual grinding method using mortar and pestle was compared to different bead beating homogenizers using different beads [1].

MATERIAL

- Precellys 24 homogenizer vs Mortar and pestle.
- Precellys kits: CK14, CK28, VK01, VK05 (2mL vials).
- Sample: ~150 mg of mycelium of *A. flavus* NRRL 3357.
- Lysis buffer: 750 µL, either TRIzol or RLT buffer (provided with Qiagen RNeasy plant mini kit) supplemented with β-mercaptoethanol.

PROTOCOL

- Precellys 24: 6500 rpm, 2x25 secs (5 sec break).
- The lysate was centrifuged at +4°C for 5-10 minutes at 16000 g for an initial homogenization.
- RNA purification: Qiagen RNeasy plant mini kit, QIAcube® robotic workstation.

[1] G.M. Leite, N. Magan, A. Medina. 2012. Comparison of different bead-beating RNA extraction strategies: an optimized method for filamentous fungi. *Journal of Microbiological Methods*, 88, 413-418.



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CONCLUSION

The use of the **Precellys 24** with the prefilled **Precellys lysing kit VK05** significantly improved the concentration and the quality of the total RNA extracted from **mycotoxigenic filamentous fungi** while reducing the required amount of mycelium. The ease of use and the time reduction aided in the molecular studies of **mycotoxigenic filamentous fungi**.

RESULTS

Significant differences were found between the total RNA amounts isolated using the Precellys 24 homogenizer and the manual system (p-value=0.0072). Furthermore the use of glass beads resulted in a yield around 3 times higher than using the traditional method of the mortar and pestle (Figure 1). The high quality of the RNA as also achieved as an example the electropherogram of a high quality RNA sample extracted using the 0.5 mm glass beads (Figure 2).

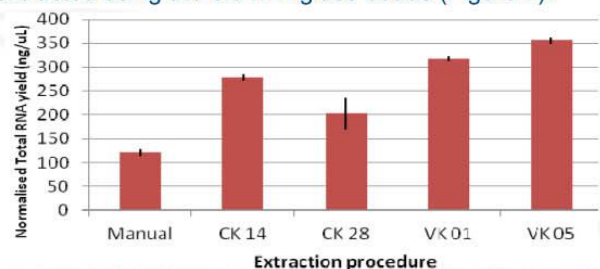


Figure 1: Total RNA yield average per 100 mg of initial biomass and standard deviation comparing different beads with the manual method. Key to treatments: CK – Zirconium Oxide, VK – Glass; followed by the bead size code 01 – 0.1 mm, 05 – 0.5 mm, 14 – 1.4 mm, 28 – 2.8 mm.

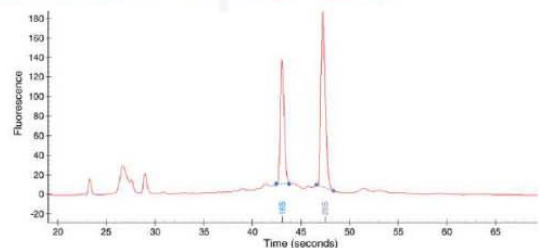


Figure 2: Electropherogram of a good quality RNA sample extracted using Precellys at room temperature. The RQI of this sample is 9.7.



For more details, please contact precellys@bertin.fr

