

Proposal for oral presentation

Dynamics and Biodiversity of microorganisms (fungi and yeast) by PCR-DGGE, with the objective of understanding Ochratoxin production on coffee beans

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Ochratoxin A (OTA) is a secondary metabolite produced by various filamentous fungi, is deemed to have nephrotoxic, immunotoxic, teratogenic and carcinogenic effects (1). In tropical zones, OTA is mainly produced in coffee beans by three *Aspergillus* species: *A. carbonarius*, *A. niger* section Nigri and *A. ochraceus* section Circumdati. In temperate zones *Penicillium verrucosum* and *P. nordicum* are known to synthesize OTA in food commodities (2, 3).

The OTA content in coffee was shown to be closely link to harvesting conditions, post-harvest processing conditions and especially dry processing, storage and transportation conditions (4, 5, 6). In order to understand the OTA contamination process in foodstuffs, assays using PCR-DGGE (Polymerase Chain Reaction - Denaturing Gradient Gel Electrophoresis) were carried. PCR-DGGE is a rapid molecular technique that was developed to monitor the dynamics of microbial populations (fungi, yeast and bacteria) by example in coffee mycoflora. In this work, the PCR-DGGE stages were optimized: extraction and amplification, repeatability and sensibility methodology applied to fungi were tested.

Methodology

Sensibility test was applied to selected *Aspergillus* species cultivated on Potato Dextrose Agar medium (AES, Combourg, France) for 5 days at 25°C. The fungal spores were collected from plates in aseptic conditions using sterile distilled water with 0.1% Tween 80 solution. The suspensions of fungal spores were quantified with Thomas cell and dilute in peptone water from 10⁷ to 10² spores/mL

DNA extraction method, according to El Sheikha *et al.* (7) different steps of extraction were used (mechanical/enzymatic/chemical). Additionally, the successful application of a eukaryotic universal primer for PCR permitted to amplify and identify many fungi species in one PCR step. The PCR products were analyzed by DGGE by using a Bio-Rad DcodeTM universal mutation detection system (Bio-Rad Laboratories, USA) using the procedure described by El Sheikha *et al.* (8).

Conclusion

The dynamics of fungus populations linked to OTA production, as well as post-harvest phytopathogens, could be studied by PCR-DGGE genetic fingerprinting. The advantages of this method are it efficiency on all microbial species (fungi, yeast and bacteria) and on the possibility of analysing a wide number of samples (30 samples) in a unique batch.

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