

# CO-CULTIVATION OF *Aspergillus nidulans* STRAINS OVER-EXPRESSING HEMICELLULASES FOR ENZYMATIC COCKTAIL PRODUCTION

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In today's world, there is an increasing trend towards the use of renewable, cheap and readily available biomass in the production of a wide variety of fine and bulk chemicals in different bioprocesses. Biorefineries utilize the enzymes of microbial cells to convert biomass into target products. The *Aspergillus niveus* genomic DNA was sequenced by pyrosequencing method and it was identified around 180 target genes encoding to cell wall-degrading enzymes. It was amplified four genes from *Aspergillus niveus* genomic DNA by PCR reactions encoding for  $\alpha$ -L-arabinofuranosidase (ABFase), endo-xylanase (XAN),  $\alpha$ -1,5-arabinanase (Arab43) and xyloendoglucanase (XEG). These genes were cloned into pExPYR vector and transformed in *A. nidulans* through protoplasts and polyethylene DNA fusion. The transformants were selected based on their zeocin-resistance and loss of the uracil and uridine auxotrophy. After that, these strains were cultivated in defined liquid minimum medium added of 5% maltose to pExPYR promoter induction followed by SDS-PAGE protein profile analysis. All described enzymes were over-expressed. Intending the production of an enzymatic cocktail, spores of the four strains over-expressing ABFase, XAN, Arab43 and XEG were inoculated in 5% maltose minimum medium. SDS-PAGE analysis showed a simultaneous production of these four enzymes. The proportion and specific activity of ABFase, XAN, Arab43 and XEG was: 4.23%/5.4 U/mg; 30%/260 U/mg; 40%/67.3 U/mg; 25.3%/33.3 U/mg, respectively. The fungi co-cultivation for simultaneous enzymes expression is an important approach to reduce the enzymatic production cost.

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