

Cloning and characterization of p450-4 and p450-1 gene involved in GA biosynthesis from *Fusarium proliferatum* KGL0401

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Objectives

GAs are secondary metabolites of the fungus *Gibberella fujikuroi*, that control such diverse developmental processes as seed germination, stem elongation, leaf expansion, trichome development, and flower and fruit development. *Fusarium proliferatum* KGL0401 was isolated from *Physalis alkekengi* var. *francheti* plant roots and it showed higher GAs productivity than wild-type *G. fujikuroi*. It was known that the seven genes (*des*, *p450-4*, *p450-1*, *p450-2*, *ggs2*, *cps/ks*, *p450-3*) of GA biosynthesis were clustered in the fungus *G. fujikuroi*. Here we present the structure of *p450-4* and *p450-1* involved in GA biosynthesis from fungus *F. proliferatum* KGL0401.

Materials and Methods

○ Materials

Fusarium proliferatum KGL0401 was isolated from *Physalis alkekengi* var. *francheti* plant roots and it showed higher GAs productivity than wild-type *G. fujikuroi*.

○ Methods

For DNA isolation the *F. proliferatum* KGL0401 were grown in 40 ml Czapek's liquid medium for 7 days at 30°C on rotary shaker set at 160 rpm. The mycelia were harvested by filtration and lyophilized for 48h. Genomic DNA was isolated from lyophilized mycelium by CTAB buffer.

Genomic DNA of *F. proliferatum* KGL0401 was used as templates for amplification of the *p450-4* and *p450-1* gene. The specific primers, which were synthesized by Bioneer Co.

Results and Discussion

p450-4 (ent-kaurene oxidase) and *p450-1* (GA14 synthase) genes were cloned and determined from a fungus *F. proliferatum* KGL0401. The deduced amino acid sequences of *p450-4* and *p450-1* showed 91% and 81% similarities with those of *G. fujikuroi*.

