

Plant Molecular Biology Update

Sequence analysis of a chalcone isomerase cDNA of *Phaseolus vulgaris* L.

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E F L H F P A V V T S P A T A K T Y F L
GGCGGCGCAGGGGAGAGAGGATTGACGATTGAGGGGAAATTCATAAAGTTCACGGCCATC 120
G G A G E R G L T I E G K F I K F T A I
GGAGTATACTTGGAGGATAAAGCGGTGGCGTCACTCGCAACAAAGTGAAGGGTAAGCCA 180
G V Y L E D K A V A S L A T K W K G K P
TCGGAAGAGTTGATCAACACTCTTGACTTCTACAGAGATATCATTTCAGGACCCCTTTGAA 240
S E E L I N T L D F Y R D I I S G P F E
AAGCTAATAAGAGGTTTGAAGATTCTGCAATTGAGTGGGACAGAATACTCGAGGAAGGTG 300
K L I R G S K I L Q L S G T E Y S R K V
ATGGA AAACTGCGTGGCACACTTGAAGTCTGTTGGGACATATGGTGATGCTGAAGCCAAA 360
M E N C V A H L K S V G T Y G D A E A K
GGCATTGAAGAGTTTGCAGAAGCCTTCAAGAAAGTGAATTTTCCACCTGGTGCCTCTGTT 420
G I E E F A E A F K K V N F P P G A S V
TTCTACCGACAATCACCTGATGGAATCTTGGGGCTTAGTTTCTCTGAAGATGCAACGATA 480
F Y R Q S P D G I L G L S F S E D A T I
CCGGGAGAAGAGGCTGTAGTTATAGAGAACAAGGCTGTATCAGCTGCAGTATTGGAGACT 540
P G E E A V V I E N K A V S A A V L E T
ATGATCGGAGAACATGCTGTCTCCCTGACTTGAAACGTAGTTTGGCTTACGATTGCTG 600
M I G E H A V S P D L K R S L A S R L L
CGGTATTGAATGGCGGCATTATAGTCTGAAAATTGAGGAACACATGAGTTGAGGAAAATT 660
R Y
AGTGCATGAACCTGAGTTTTTCAAAATATTATTGTTTTTCTCTTTTCTTCTCACTAAAGT 720
TGTTAAACTTACTCATGTATTTCTACTTGTTTATCAAATAAATAAATAAATCTGAGTTT 780
GAGATTTTCTTAAAAA AAAAAAAAAA
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Fig. 1. The nucleotide and derived amino acid sequences of the bean CHI cDNA clone in pCHI1. Potential polyadenylation signals are underlined.

The nucleotide sequence data reported will appear in the EMBL, GenBank and DDBJ Nucleotide Sequence Databases under the accession number X16470.

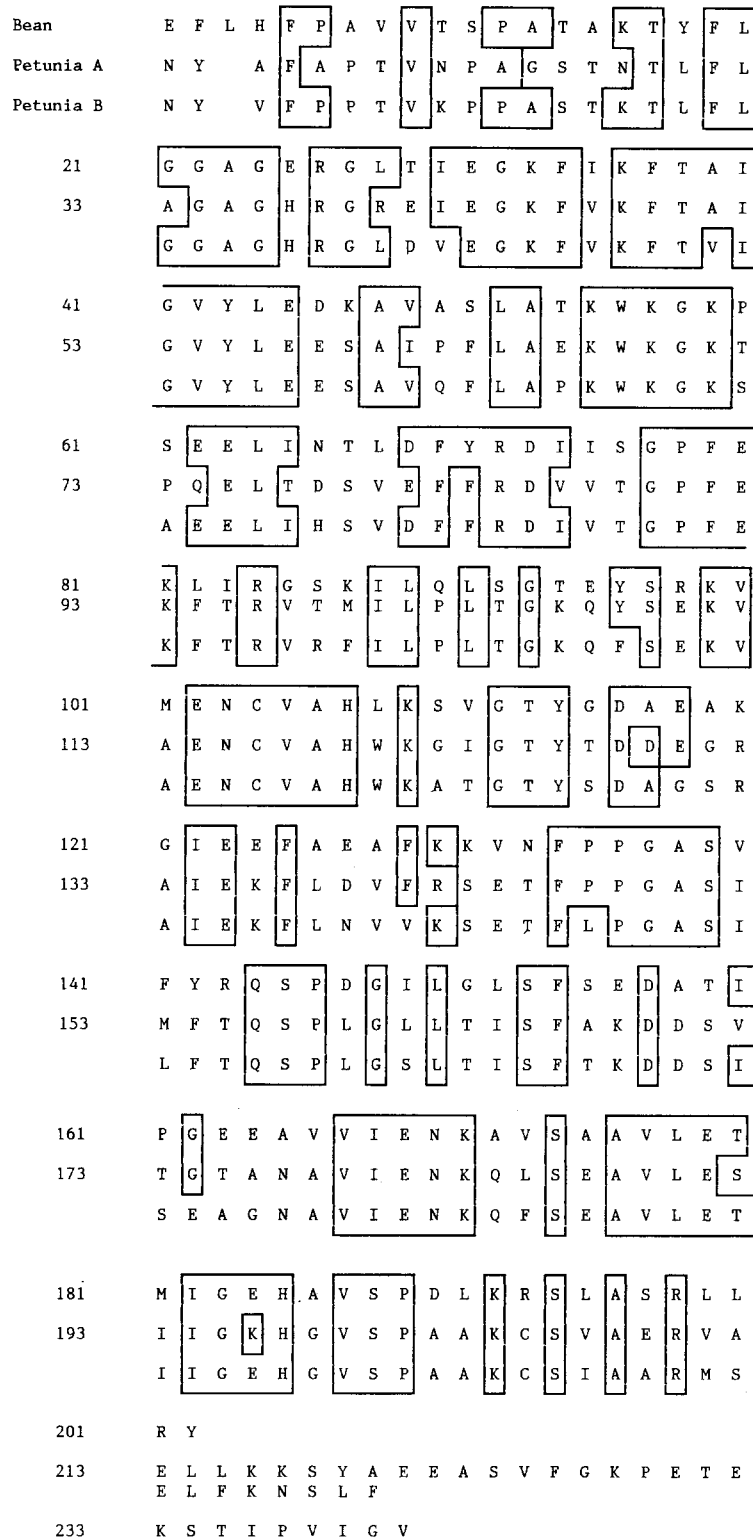


Fig. 2. Comparison of the derived amino acid sequences of chalcone isomerases from bean (CHI1) and *Petunia hybrida* (CHIA and CHIB).

Chalcone isomerase (CHI, E.C. 5.5.1.6) catalyzes the stereospecific isomerization of chalcones to their corresponding minus (-) flavanones, a key reaction in the biosynthesis of flavonoid pigments and, in the Leguminosae, in the formation of antimicrobial isoflavonoids [1]. A λ gt 11 expression library was constructed from mRNA from bean (*Phaseolus vulgaris*) suspension cells treated with fungal elicitor [2]. This treatment results in increased levels of CHI protein and enzyme activity [3]. The library was screened with an antibody raised against purified CHI from the bean cultures [3], and two putative CHI clones were isolated [2]. The approximately 800 bp insert in clone pCHI1 was identified as encoding CHI based on the ability of the insert sequence to hybrid select an mRNA encoding a 27 kDa polypeptide recognized by the anti-(bean CHI) serum [2]. Use of this insert in northern blot experiments revealed induction of bean CHI transcripts in elicitor-treated cell suspension cultures, wounded hypocotyls, and hypocotyls infected with compatible and incompatible races of the fungal pathogen *Colletotrichum lindemuthianum* [2].

We now report the nucleotide and deduced amino acid sequence of the insert in clone pCHI1. The insert sequenced consisted of 808 nucleotides, comprising 185 nucleotides of 3'-untranslated region (with three potential polyadenylation signals), a short poly(A) tail, and 606 nucleotides of coding sequence (Fig. 1). Comparison with other CHI sequences [4] and with the M_r of the bean protein [3] suggests that the clone is lacking approximately 36 nucleotides of the 5' coding sequence. pCHI1 was, however, the longest CHI clone obtained from the λ gt 11 library. At the amino acid level, the bean CHI is 59% homologous to the CHIA and CHIB polypeptides from *Petunia hybrida* [4] (Fig. 2). The regions of homology occur in short blocks throughout the sequence. This might explain the lack of any strong antigenic cross-reactivity between the CHIs from bean and *Petunia* [5]. Hydropathy plots for the bean and *Petunia* enzymes are, however, extremely similar (data not shown).

CHI from bean catalyzes the isomerization of

both 2',4,4'-trihydroxychalcone and 2',4,4',6'-tetrahydroxychalcone to their corresponding flavanones, whereas the *Petunia* enzyme is inactive with the trihydroxychalcone [5]. These specificities reflect the A-ring substitution patterns of the flavonoid derivatives found in the two plant species [1]. The phytoalexins of *Petunia* are not isoflavonoids, so in this species CHI functions specifically in the formation of flavonoid pigments from 2',4,4',6'-tetrahydroxychalcone. The trihydroxychalcone A-ring substitution pattern is common to most of the antimicrobial isoflavonoids of the Leguminosae. Work is currently in progress, using molecular modeling and site-directed mutagenesis, to understand the molecular basis of the different substrate specificities of the CHIs from bean and *Petunia*.

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