





## **Biosafety Clearing-House (BCH)**

**GENETIC ELEMENT (GENE)** 

BCH-GENE-SCBD-100267-2

LAST UPDATED: 16 JUL 2012

#### **General information**

Name of genetic element

delta(12)-fatty acid dehydrogenase

ΕN

Abbreviation

CS-gm-fad2-1

EN

Category

Protein coding sequence

Is this genetic element a synthetic molecule?

No

#### **Donor organism**

Donor organism(s)

BCH-ORGA-SCBD-10453-6 ORGANISM | GLYCINE MAX (SOYBEAN, SOYA BEAN, SOYA, SOYBN) | Crops

### Characteristics of the protein coding sequence

Name of the protein expressed by the coding sequence

delta(12)-fatty acid dehydrogenase

ΕN

Biological function of the protein

The gm-fad2-1 is a fragment of the omega-6 desaturase gene 1 (FAD2-1) from Glycine max (soybean) that corresponds to approximately 40% of the middle portion of the coding region of FAD2-1 (597 bp).

ΕN

gm-fad2-1 itself does not code for a functional protein, but transcription of this gene fragment transgenic soybean seeds acts to suppress transcription of endogenous omega-6 desaturase, resulting in the high oleic phenotype.

Related trait(s) or use(s) in biotechnology

Changes in quality and/or metabolite content Lipid and fatty acids

#### **Additional Information**

Other relevant website addresses and/or attached documents

? GM Crop Database - CERA ( <code>English</code> )

 $\ref{eq:continuous}$  Delta12-fatty acid dehydrogenase - Wikipedia (  $\it English$  )

BCH-GENE-SCBD-100267-2

# **Further Information**

Questions about the Cartagena Protocol on Biosafety or the operation of the Biosafety Clearing-House may be directed to the Secretariat of the Convention on Biological Diversity.

Secretariat of the Convention on Biological Diversity

413 rue Saint-Jacques, suite 800 Montreal, Québec, H2Y 1N9 Canada

Fax: +1 514 288-6588 Email: secretariat@cbd.int