





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

LIVING MODIFIED ORGANISM (LMO)

BCH-LMO-SCBD-46334-8

### ? Decisions on the LMO ? Risk Assessments

LAST UPDATED: 28 JAN 2013

## **Living Modified Organism identity**

The image below identifies the LMO through its unique identifier, trade name and a link to this page of the BCH. Click on it to download a larger image on your computer. For help on how to use it go to the LMO quick-links page.

https://bch.cbd.int/database/record?documentID=46334



BCS-GHØØ2-5 GlyTol™ Cotton GHB614



Read barcode or type above URL into internet browser to access information on this LMO in the Biosafety Clearing-House @ SCBD 2012

Name

GlyTol™ Cotton GHB614

ΕN

Transformation event

**GHB614** 

Unique identifier

BCS-GHØØ2-5

Developer(s)

- PERSON: BAYER CROPSCIENCE | BCH-CON-CA-7708-1

**PERSON** 

Bayer CropScience

#### Description

GlyTol cotton event GHB614 contains a stably integrated gene 2mepsps. The 2mepsps gene was generated by introducing mutations into the wild-type epsps (wt epsps) gene from maize, leading to a double mutant EPSPS protein with two amino acid substitutions (2mEPSPS). This modification confers the protein a decreased binding affinity for glyphosate, allowing it to maintain sufficient enzymatic activity in the presence of the herbicide. Therefore, the plants bearing this gene are tolerant to glyphosate herbicides.

ΕN

#### Recipient Organism or Parental Organisms

The term "Recipient organism" refers to an organism (either already modified or non-modified) that was subjected to genetic modification, whereas "Parental organisms" refers to those that were involved in cross breeding or cell fusion.

BCH-ORGA-SCBD-12080-6 ORGANISM GOSSYPIUM HIRSUTUM (COTTON)

Crops

Point of collection or acquisition of the recipient organism or parental organisms

Variety: Coker 312 ΕN

# Characteristics of the modification process

Vector

pTEM2 ΕN

Techniques used for the modification

Agrobacterium-mediated DNA transfer

Genetic elements construct

V-LB-RHIRD 0.024 kb

P-h4a748-ARATH I-H3-ARATH 1.011 kb

0.516 kb

TP-OPT 0.372 kb

CS-epsps-MAIZE 1.337 kb

T-H4-ARATH 0.742 kb

V-RB-RHIRD 0.024 kb

#### Introduced or modified genetic element(s)

Some of these genetic elements may be present as fragments or truncated forms. Please see notes below, where applicable.

## BCH-GENE-SCBD-46333-8 5-ENOLPYRUVYLSHIKIMATE-3-PHOSPHATE SYNTHASE | (MAIZE, CORN)

Protein coding sequence | Resistance to herbicides (Glyphosate)

BCH-GENE-SCBD-101419-4 OPTIMIZED TRANSIT PEPTIDE

Transit signal

BCH-GENE-SCBD-101415-9 TI PLASMID LEFT BORDER REPEAT

Plasmid vector

BCH-GENE-SCBD-104647-3 HISTONE H4 GENE PROMOTER | (THALE CRESS)

Promoter

BCH-GENE-SCBD-104648-2 HISTONE H3 GENE II INTRON 1 | (THALE CRESS)

Intron

BCH-GENE-SCBD-104646-4 HISTONE H4 GENE 3' UTR | (THALE CRESS)

Terminator

BCH-GENE-SCBD-101416-6 TI PLASMID RIGHT BORDER REPEAT

Plasmid vector

Notes regarding the genetic elements present in this LMO

Southern blot analysis indicated that a single insertion of the complete T-DNA as constructed occurred in the GHB614 line. There was no indication that any portion of the vector backbone also integrated into the GHB614 line.

ΕN

## **LMO** characteristics

Modified traits

Resistance to herbicides
Glyphosate

Common use(s) of the LMO

Food

Feed

Fiber/textile

# **Detection method(s)**

External link(s)

? BCS-GHØØ2-5 - EU Reference Laboratory for GM Food and Feed (EURL-GMFF) ( English )

? BCS-GHØØ2-5 - CropLife International Detection Methods Database ( English )

#### **Additional Information**

Other relevant website addresses and/or attached documents

? Bayer CropScience: Petition for Determination of Nonregulated Status for Glyphosate-Tolerant cotton.pdf ( English )

? GMO Compass ( English )

BCH-LMO-SCBD-46334-8

# **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

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