



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

#### LIVING MODIFIED ORGANISM (LMO)

BCH-LMO-SCBD-115143-2

#### ? Decisions on the LMO ? Risk Assessments

LAST UPDATED: 21 JAN 2022

#### 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=115143

Cassava brown streak disease-resistant cassava

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

Name

Cassava brown streak disease-resistant cassava

Transformation event

D5001-985013

Developer(s)

#### - ORGANIZATION: NATIONAL ROOT CROPS RESEARCH INSTITUTE || BCH-CON-SCBD-115138-1

#### ORGANIZATION

National Root Crops Research Institute Academic or research institute Km 8 Ikot Ekpene Road, PMB 7006 Umuahia, Abia Nigeria Phone: +234 (0)8168983790, +2349035068714 Email: info@nrcri.gov.ng, Ihuomaumezurumba@yahoo.com Website: https://nrcri.gov.ng/

#### Description

Cassava (*Manihot esculenta* Crantz) was modified for RNA interference-mediated resistance to Cassava Brown Streak Disease. The modified cassava expresses a hairpin RNA (hpRNA) cassette that contains portions of the coat proteins of *Cassava brown streak virus* and *Ugandan cassava brown streak virus*, the causal agents of Cassava Brown Streak Disease. Transcription of the hpRNA form this cassette guides host cell machinery for targeted degradation of infecting viral transcripts.

EN



ΕN

# A selectable marker, *Escherichia coli* neomycin phosphotransferase II, was also included for selection of transformants using kanamycin.

#### 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-112539-1 ORGANISM | MANIHOT ESCULENTA (CASSAVA, BRAZILIAN ARROWROOT, YUCA, MANIOC, MANDIOCA, MANES)

Crops

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

Cassava cultivar: TMS 98/0505

ΕN

EN

Related LMO(s)

BCH-LMO-SCBD-115144-1Viral resistant cassava with increased levels of zinc and iron | Dr IhuomaOkwuonu | Changes in quality and/or metabolite content, Increased iron levels, Increased zinclevels, Resistance to antibiotics (Kanamycin), Resistance to CBSV, Resistance to diseases and pests(Viruses), Resistance to UCBSV, Selectable marker genes and reporter genesBCH-LMO-SCBD-115140-1Viral resistant cassava with increased levels of zinc and iron | NationalRoot Crops Research Institute(NRCRI) | Changes in quality and/or metabolite content, Increased ironlevels, Increased zinc levels, Resistance to antibiotics (Kanamycin), Resistance to CBSV, Resistance to CBSV, Resistanceto diseases and pests (Viruses), Resistance to UCBSV, Selectable marker genes and reporter genes

#### Characteristics of the modification process

Vector

p5001

Techniques used for the modification

Agrobacterium-mediated DNA transfer

Genetic elements construct

	T-nos-RHIR 0.266 kb	CS-cp_RNAi-C 0.897 kb	CS-cp_RNAi-UC 0.899 kb	I-3_pdk-FLA. 0.781 kb	CS-cp_RNAi-UCBSV 0.899 kb	CS-cp_RNAi-CBSV 0.897 kb
P 0	-CsVMV .515 kb					
	T-nos-RHIR 0.249 kb	CS-nptII-EC 0.795 k	P-e35S-CaMV 0.781 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-101900-6 CSVMV PROMOTER | CASSAVA VEIN MOSAIC VIRUS (CASSAVA VEIN MOSAIC VIRUS, CVMV, CSVMV)

Promoter

BCH-GENE-SCBD-115110-2 COAT PROTEIN | UGANDAN CASSAVA BROWN STREAK VIRUS (UCBSV)
Double-stranded RNA

BCH-GENE-SCBD-115108-2 COAT PROTEIN | CASSAVA BROWN STREAK VIRUS (CASSAVA BROWN STREAK VIRUS, CBSV)

Double-stranded RNA

BCH-GENE-SCBD-103123-6 PYRUVATE ORTHOPHOSPHATE DIKINASE, INTRON 3 | FLAVERIA TRINERVIA (CLUSTERED YELLOWTOPS, SPEEDYWEED, FLAVERIA, YELLOW TWINSTEM)

Intron

BCH-GENE-SCBD-100269-8 NOPALINE SYNTHASE GENE TERMINATOR

Terminator

BCH-GENE-SCBD-100366-6 CAMV ENHANCED 35S PROMOTER
Promoter

BCH-GENE-SCBD-15001-5 NEOMYCIN PHOSPHOTRANSFERASE II | (BACTERIA)

Protein coding sequence | Resistance to antibiotics (Kanamycin)

Notes regarding the genetic elements present in this LMO

#### **RNA interference cassette:**

Transcription of the cassette produces a hairpin RNA (hpRNA). RNA polymerase is recruited to the Cassava vein mosaic virus promoter and then transcribes: two segments of *Cassava brown streak virus* (CBSV) and the *Ugandan cassava brown streak virus* (UCBSV) coat protein (sense orientation), the *Flaveria trinervia* pyruvate orthophosphate dikinase intron 3 (*pdk*), and CBSV and UCBSV coat protein (anti-sense orientation). Transcription stops at the *Agrobacterium tumefaciens* nopaline synthase (*nos*) terminator. The produced RNA then can form a double stranded segment due to the inter-strand homology between the CBSV and UCBSV coat protein segments in inverted orientations and the flexible linker/loop, *pdk*. The double stranded segment of the hpRNA is sufficient to elicit an RNA interference (RNAi) response. The host cell machinery then processes the hpRNA into small interfering RNA, which is used to guide the targeted degradation of RNA molecules with sequence homology. In this case, infecting viral transcripts are degraded by the host cell.

Please note all genetic elements were in the anti-sense orientation.

#### LMO characteristics

Modified trait	ts				
Desistence to discourse and packs					
Resistance	ance to diseases and pests				
	Viruses				
Resistance	ance to antibiotics				
	Kanamycin				
Other					
	Resistance to CBSV				
	Resistance to UCBSV				

Food Research

#### **Detection method(s)**

Additional Information

Due to the processing of the hpRNA by the host cell DICER protein, it is expected that small interfering RNA 21 to 24 base pairs in length will be present instead. Additionally, the cellular processing means that no proteins are expected to be produced from the hpRNA.

#### **Additional Information**

Additional Information

Please note that several lines have been created using the transformation vector p5001. The following transformation events are expected to contain similar genetics:

D5001-985013, D5001-985019, D5001-985022, D5001-985024, D5001-985025, D5001-985026, D5001-985028, D5001-985029, D5001-985030, D5001-985031, D5001-985033 and D5001-985034.

The transformation event is thus a placeholder until further information is uncover and/or a line has been selected for commercialization.

Other relevant website addresses and/or attached documents

? Virus-Derived Stacked RNAi Construct Confers Robust Resistance to Cassava Brown Streak Disease.pdf (*English*)

#### BCH-LMO-SCBD-115143-2

## **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