





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

# LIVING MODIFIED ORGANISM (LMO)

BCH-LMO-SCBD-115602-1

### ? Decisions on the LMO ? Risk Assessments

LAST UPDATED: 11 JUN 2020

# **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=115602



Banana modified for Xanthomonas wilt disease resistance



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

#### Name

### Banana modified for Xanthomonas wilt disease resistance

ΕN

Transformation event

#### **HRAP**

### Developer(s)

- PERSON: DR. LEENA TRIPATHI | BCH-CON-SCBD-115601-2

### **PERSON**

Dr. Leena Tripathi Principal Scientist Nairohi

Nairobi Kenya

Phone: +254 208632900 Email: I.tripathi@cgiar.org

RELATED ORGANIZATION

### Description

The banana was modified for resistance to Xanthomonas wilt disease, caused by *Xanthomonas campestris* pv. *musacearum* through the expression of *Capsicum annuum* hypersensitive response-assisting protein. The modified banana plants demonstrate more a robust hypersensitive response upon infection with Gram negative bacteria, such as *X. campestris*, and thus resist disease related wilting. The modified banana also contains a selectable marker, *Ecscherichia coli* neomycin phosphotransferase II, for kanamycin selection

ΕN

### during transformation.

# 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-100304-3 ORGANISM | MUSA SP. (BANANA, PLANTAIN)

Crops

#### Related LMO(s)

BCH-LMO-SCBD-115604-1 | Banana resistant to Xanthomonas wilt disease | Dr. Leena Tripathi

Resistance to antibiotics - Kanamycin Resistance to diseases and pests - Bacteria

Show detection method(s)

BCH-LMO-SCBD-115603-1 | Banana modified for Xanthomonas wilt disease resistance | Dr. Leena

Tripathi Resistance to antibiotics - Kanamycin Resistance to diseases and pests - Bacteria

Show detection method(s)

# Characteristics of the modification process

Vector

pBI-HRAP EN

Techniques used for the modification

# Agrobacterium-mediated DNA transfer

#### Genetic elements construct

V-RB-RHIRD	P-nos-RHIRD	CS-nptII-ECOLX	T-nos-RHIRD
0.000 kb	0.180 kb	0.795 kb	0.253 kb
P-35S-CaMV	CS-HRAP-CAPAN	T-nos-RHIRD	V-LB-RHIRD
0.346 kb	0.834 kb	0.253 kb	0.000 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-101416-6 TI PLASMID RIGHT BORDER REPEAT

Plasmid vector

BCH-GENE-SCBD-100270-6 NOPALINE SYNTHASE GENE PROMOTER

Promoter

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

Protein coding sequence | Resistance to antibiotics (Kanamycin)

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

Terminator

BCH-GENE-SCBD-100287-7 CAMV 35S PROMOTER

Promoter

BCH-GENE-SCBD-113355-1 HYPERSENSITIVE RESPONSE ASSISTING PROTEIN GENE | CAPSICUM

ANNUUM (BELL PEPPER, SWEET PEPPER, CHILI PEPPER, CAPAN)

Protein coding sequence | Resistance to diseases and pests

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

Plasmid vector

Notes regarding the genetic elements present in this LMO

# Gene expression

Transcription of *Escherichia coli* neomycin phosphotransferase II is under control of the *Agrobacterium tumefaciens* nopaline synthase (*nos*) promoter and terminator.

Transcription of *Capsicum annuum* hypersensitive response assisting protein (*hrap*) is under control of the Cauliflower Mosaic Virus 35S promoter and the *nos* terminator. High levels of transcription are expected due to the constitutive nature of the CaMV promoter.

ΕN

#### Notes:

- The transformation vector was derived from the pBI121 vector using a BamHI and SacI restriction digest.
- The coding sequence of hrap was derived from the cDNA clone of the mRNA.

### **LMO** characteristics

Modified traits

Resistance to diseases and pests

Bacteria

Resistance to antibiotics

Kanamycin

Selectable marker genes and reporter genes

Other

Resistance to Xanthomonas sp.

Common use(s) of the LMO

Food

Research

# **Detection method(s)**

External link(s)

? Expression of sweet pepper Hrap gene in banana enhances resistance to Xanthomonas campestris pv musacearum.pdf ( English )

Additional Information

Some detection methods can be found in the scientific article detailing the creation of this banana event.

ΕN

# **Additional Information**

Other relevant website addresses and/or attached documents

- $\ref{eq:continuous}$  Field trial of Xanthomonas wilt disease-resistant bananas in East Africa.pdf ( <code>English</code> )
- $\ref{eq:continuous}$  Expression of sweet pepper Hrap gene in banana enhances resistance to Xanthomonas campestris pv musacearum.pdf ( English )
- ? SnapGene: pBI121 (empty vector) ( English )

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# **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