



European Commission - Directorate General Joint Research Centre

## Institute for Health and Consumer Protection Molecular Biology and Genomics Unit

### Standard Operating Procedure DNA Extraction from honey and pollen – CTAB

Revision	Author	Reviewer	QS Agreement	Approver
1	I. Matetovici	M. Van den Bulcke ...	S. Cordeil	M. Mazzara
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## Molecular Biology and Genomics *Standard Operating Procedure*

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## **1 Object**

This method is used to extract genomic DNA from maize, soy, rapeseed pollen present in honey.

## **2 Applicability domains**

The method for DNA extraction described below is suitable for the isolation of genomic DNA from maize, soy, rapeseed pollen in honey. Application of the method to other substances may require adaptation and possible further specific optimisation.

## **3 Reference documents**

Folloni S., Kagkli D.M., Rajcevic B. Guimarães N., Valicente F., Van Droogenbroeck B., Van den Eede G. Van den Bulcke M. Detection of airborne genetically modified maize pollen by real-time PCR (*in press*)

## **4. Abbreviations used**

CTAB: Cetyltrimethylammonium Bromide

DNA: Deoxyribonucleic acid

EDTA: Ethylenediaminetetraacetic acid

RNase: Ribonuclease

Tris: Tris[hydroxymethyl] aminomethane

## **5. Equipment**

Water bath or heating block

12µM membrane filter (cellulose nitrate)

Vacuum filtration system (Sartorius)

Microcentrifuge (Eppendorf or equivalent)

Micropipettes

Vortexer

Microcentrifuge tubes

Rack for reaction tube

Vinyl or later gloves

Vacuum device

Mini-shaker IKA (or equivalent)

1 mm glass beads



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#### **NOTE:**

Plastic ware has to be sterile and free of DNases, RNases and nucleic acids. Filter pipette tips protected against aerosol should be used.

### **6. Reagents**

2% CTAB- buffer

Chloroform (99% at least)

Isopropanol (99, 7 % at least)

Ethanol (96% at least)

Ethanol-solution 70% (v/v)

Proteinase K solution 20 mg/ml

RNase A solution 0.5 mg/ml

Sterile molecular biology grade water

### **7. Procedure**

#### **PRELIMINARY REMARKS:**

- The procedure requires sterile conditions. Contamination may be avoided during sample preparation by using single-use equipment, decontamination solutions and by avoiding the formation of dust.
- During all steps of DNA preparation, wear gloves and change gloves regularly.
- Avoid extensive vortexing of genomic DNA.

#### **Step 1: Honey solubilisation and pollen fraction collection**

- 1) Heat the honey jar at 40-50 °C until the matrix becomes fluid.
- 2) Weight the desired amount of honey to a proper recipient (e.g. a 50 ml Falcon tube when analysing 10g of honey)
- 3) Add 4 volumes of preheated (40-50 °C) sterile deionised water.
- 3) Put the samples on an agitator and rotate them until the honey is completely dissolved.



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4) After the honey is completely dissolved filter the wet solution on a 12 $\mu$ M filter using a Vacuum filtration system (Sartorius) or equivalent.

5) Transfer the filters into an appropriate recipient (e.g. a 15 ml Falcon tube).

#### Notes:

- When using large amounts of honey, the 12 $\mu$ M filter may become saturated with material and filtering hampered. Take the filter out of the filter unit, transfer to the Falcon tube and replace by a new 12 $\mu$ M filter and continue the filtration. This handling can be repeated until the complete honey solution has been filtered.
- Honey contains in general low amounts of pollen (in the range of 1000 pollen grains/gram honey). A suitable amount for analysis may be 10-50g, although larger amounts may be required (e.g. 500g).
- Filters with pollen material can be stored for short periods at 4°C. Storage for longer periods is to be avoided but if the case, air-dried filters can be stored in a Falcon tube at minus 20 °C

### **Step II: Isolation and Purification of genomic DNA of collected pollen fraction**

- 1) Take the filter(s) in the Falcon tube
- 2) Add 1 ml of molecular biology grade water
- 3) Add 1 volume of glass beads (1 mm glass beads)
- 4) Add 2 ml of 2% CTAB-buffer
- 5) Shake at maximum strength using a mini-shaker IKA (or equivalent) to disrupt the pollen grains; first shake for about 1 min, and then allow settling for about 30 sec on ice and finally shake again for 2 min
- 6) Add 50  $\mu$ l Proteinase K and incubate with agitation at least 2 hours (or over night) at 60° C
- 7) Add 1  $\mu$ l of RNase and incubate 30 min at room temperature
- 8) Centrifuge for 15 min at 5000  $\times$  g
- 9) Transfer the supernatant in 1 ml fractions to a 2 ml Eppendorf tube and add 900  $\mu$ l of chloroform (or repeat in different tubes until the supernatant is completely collected)



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- 10) Mix for 30 sec and centrifuge at 12000  $\times$  g for 10 min
- 11) Transfer the aqueous phase to a new tube and add 0.8 volume of Isopropanol
- 12) Incubate for 30 min at room temp
- 13) Centrifuge at 12000  $\times$  g for 10 min
- 14) Discard the supernatant
- 15) Wash the pellet with 1 ml of ethanol 70%
- 16) Mix by inverting, centrifuge at 12000  $\times$  g for 5 min
- 17) Discard the supernatant and repeat the washing step
- 18) Dry the pellet (room temp or concentrator)
- 20) Re-dissolve the DNA in 60  $\mu$ l sterile molecular biology grade water. Add the water in the first tube and resuspend the DNA, then take the obtained solution and added to the next tube until all the tubes are pooled.

#### **Optional**

21) Purify the DNA samples using the Genomic DNA Clean& Concentrator<sup>TM</sup> kit (Zymo Research) according with manufacturers' instructions.

### **8. Procedure Buffer preparation and required reagents**

The following reagents/buffers/solutions are required for the DNA extraction procedure:

#### **8.1 Equipment**

Weighting equipment  
pH meter  
Heating stirrer  
Autoclave  
Pipettes and micropipettes



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## **8.2 Reagents**

**CTAB:** Cetyltrimethylammonium Bromide (Ultrapure grade)

**Tris:** Tris[hydroxymethyl] aminomethane (Molecular Biology grade) or 1 M Tris-HCl pH 8.0

**EDTA:** Ethylenediaminetetraacetic acid (titration 99.0%)

Ethanol (98% at least)

NaCl (99% at least)

NaOH (98% at least, anhydrous)

Sterile molecular biology grade water

## **8.3 Formulations**

### *2% CTAB-buffer*

20 g/l CTAB.....4 g

1.4 M NaCl.....16.4 g

0.1 M Tris\*.....3.15 g

20 mM Na<sub>2</sub>EDTA.....1.5 g

\* Alternatively 20 ml of 1M ready-to-use solution of Tris-HCl pH 8.0 can be used.

Preparation:

- add 100 ml distilled water
- adjust pH to a value of 8.0 with 1M NaOH
- fill up to 200 ml and filter sterilise
- store buffer at 4°C for max. 6 months.

### *NaOH 1 M*

Dissolve 2 g of NaOH in 50 ml sterile water

### *Ethanol-solution 70% (v/v)*

70 ml of pure ethanol are mixed with 30 ml of distilled sterile water.