# COEXISTENCE BETWEEN TRANSGENIC MON 810 MAIZE AND HIVES

Eugénia Andrade<sup>1</sup>, Maria Lopes<sup>2</sup>, Mónica Rodrigues<sup>1</sup>, Ana Chegão<sup>1</sup>, Fátima Quedas<sup>2</sup>, Amélia Lopes<sup>1</sup>

<sup>1</sup> INIAV; <sup>2</sup> IPS







#### Introduction

- Pollen is a natural constituent of honey (≤ 0.5%).
- Maize flour might be considered as an ingredient or a contaminant of honey.
- So far PCR has not been considered a helpful technique to separate pollen from contaminants.
- The aim of this study was to investigate the ability of qPCR using plasmid calibrants to provide good estimates of the transgene copy number in relation to a species specific gene copy number and look for any possible application in honey analysis.

### Material

- Spiked honey samples
  - GM-F<sub>1</sub> pollen/honey (m/m): 0.45; 0.10; 0.04; 0.01; 0.005%
  - GM-F<sub>1</sub> embryos (m/m): 0.10%
  - GM-F<sub>1</sub> flour: transgene donor female parent (m/m): 0.10%
  - GM-F1 flour: transgene donor male parent (m/m): 0.10%
- Local honey (Salvaterra)
- Imported (EU and non-EU) honey

### Methods

- DNA extraction: validated CTAB-based method (van den Bulcke et al., 2012), slightly modified.
- Calibration curves: ERM®-AD413 (for MON810) and ERM®-AD415 (for NK603) calibrants at nominal concentration of 2 x 10<sup>6</sup> copies/μL
- Quality control for PCR efficiency and positive control: ERM®-BF413f
- Real-time PCR: according to EURL GMFF, but with a plasmid calibrant certified for the copy number ratio instead of a reference material certified for its GM mass fraction.

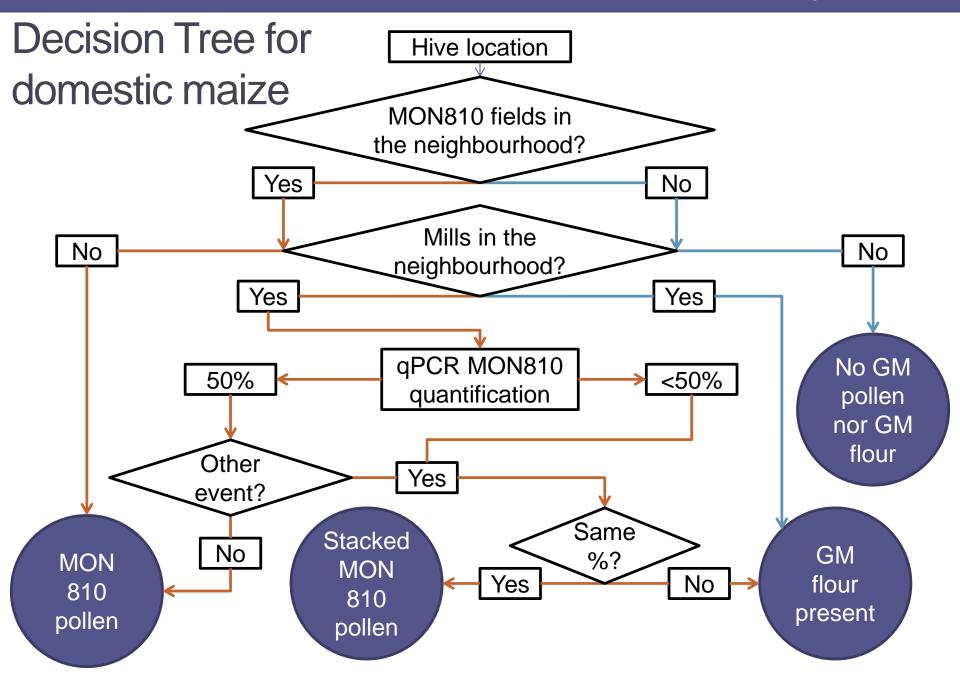
## GM copy number ratios

Matrix – Spicked samples	MON810 copy number ratio (Average ± Standard deviation)
Embryos (hemizygous)*	53.73 ± 0.09
F1 seed flour (transgene donor: ♀ parent)	59.07 ± 0.02
F1 seed flour (transgene donor: ♂ parent)	38.83 ± 0.04
Pollen (from a hemizygous plant)	52.82 ± 4.25

	MON810 copy number ratio (Average ± Standard deviation)
Salvaterra honey	52.63
Flowers honey**	24.02

<sup>\*</sup> Model for F<sub>2</sub> flour

<sup>\*\*</sup> Also positive for GM oilseed rape and NK603



### Conclusions

- The combination of qPCR and plasmid calibrants allowed to differentiate different maize tissues ploidies.
- This study confirms that PCR is not a helpful technique to separate by itself pollen from flour in honey.
- However, when dealing with single origin honey, it might be coupled with other information to determine the most likely sources of GM presence.

