## Cell Wall Modifying Gene expression during banana fruit ripening and in relationship with finger drop



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#### **Introduction and Objectives**

- Finger drop is one of the main features closely associated with the banana ripening process.
- This process implies physiological and biochemical changes leading to peel softening and weakening, thus causing individual fruit in a hand to separate from the crown (see below).

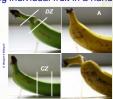
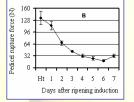


Fig 1: Cavendish banana fruit taken at green and rine stages rig 1: Caverious oriental multitakent at green and hip stagets. Peel tissue of control (CZ) and drop (DZ) zones used in this study are indicated in panel A. Typical evolution of finger drop observed during fruit ripening measured throughout the pedical rupture force was indicated in panel B.



Expression of two

- Therefore the finger drop process reduces the commercial quality and economic value of bananas as hands with fingers missing or fingers without pedicels cannot be sold to consumers.
- Although this, few studies have been devoted on finger drop process and only at physicochemical and biochemical levels (Paull, 1996; Imsabai et al., 2006; Sengpook et al., 2007).
- In the prospect of improvement of banana fruit quality throughout conventional breeding and marker assisted selection, getting major or candidate gene(s) putatively involved on finger drop process is an essential step for identification of related marker (s).
- In order to identify these candidate genes, we examined here expression of Cell Wall Modifying Genes (CWMGs) in peel tissue at median area (CZ) and compared to that in the pedicel rupture area (DZ).

#### **Materials and Methods**

- Studies were performed on peel of Cavendish banana fruit ripen in air at 20°C after azethyl (95% N + 5% C2H4) treatment (1000ppm / 24 h / 20°C).
- As CWMGs, we examined those involved in metabolism of cell wall component including pectin (PME, PEL and PG) and xyloglucan (XTH) and those involved in physical properties of cell wall (EXP).
- CWMGs were isolated by RACE-PCR (XTHs). Musa sequence data mining (PME) or systematic searches in GenBank database (PEL, PG and EXP).
- CWMGs expression was performed throughout real-time quantitative PCR comparatively at control and drop zones.

PFI = nectate lyase PG= polygalacturonase PME= pectin methylesterase XTH= xyloglucan transglycosylase hydrolase EXP= expansin



pedicel rupture (Chillet et al. 2008

### **Results**

#### Expression of four pectolytic genes xyloglucan metabolism genes ····· Control zone — Drop zone Control zone → Drop zone 8000 2500 600 (cal Ht ref actin) 360 240 120 Ht, ref actin) 6400 (cal Ht, ref actin) 2000 2-dact Mapell (cal Ht, ref actin) 4800 1500 3200 1000 120 1600 500 2000 3000 30 (cal Ht, ref actin) 1600 (cal Ht, refactin) 2400 MA-EXP2 (cal Ht, refactin) 1200 18 2,4 1800 Ή 800 12 1,6 1200 8,0 600 Ht 1 2 3 4 5 6 Ht 1 2 3 4 5 6 Ht 1 2 3 4 5 6 7 Days after ripening induction Days after ripening induction Days after ripening induction

#### Expression of four cell wall loosening genes

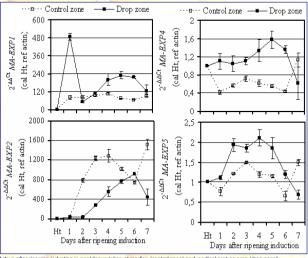


Fig 2: Expression of CWIMGs during in peel tissue and during ripening of Cavendish banana fruit. CWIMGs expression was performed daily since the harvest time (H) of from 7 days after ripening induction in peel tissue taken at median (control zone) and pedicel rupture area (drop zone)

#### **Discussions and Perspectives**

- Finger drop phenomenon implies changes in CWMGs expression involved in metabolism of cell wall components (pectin and xyloglucan) and in physical properties of cell wall.
- Theses changes occur mainly 1-4 days after ripening induction. Based on CWMGs genes, the sequence of events begins with changes in pectolitic component following and physical properties of Cell wall following by xyloglucan metabolism.
- Contrary to MaEXP2, MaPME2, MaPEL1 and 2, MaPG4, MaXTH8 and 9, MaEXP1, 4 and 5 of which mRNA accumulated highly in the drop zone appeared as good candidates.
- The next step of this study will be the identification, for breeding program, of molecular marker related from each candidate CWMG. To this end, the biodiversity among banana species will exploited to investigate the relationship between structural variability of allelic form of each CWMG - its expression - phenotype (finger drop).

#### References