

Understanding the Physiological and Biochemical Mechanisms of Graft-incompatibility

Stefano Musacchi, Associate Professor

Endowed Chair of Tree Fruit Physiology and Management

WSU Tree Fruit Research and Extension Center,
Wenatchee, WA

2015 Empire State Produces EXPO New York January 20, 2015

Why is graft-incompatibility between pear and quince a major problem in Italy?

94% of orchards in Italy are grafted on quince

Why do growers still use quince, despite graft-incompatibility?

- **Dwarfing (HDP and UHDP)**
- **Early bearing**
- **Easy to propagate**

Pear cultivars classification on the basis of graft-incompatibility level with quince

Compatible

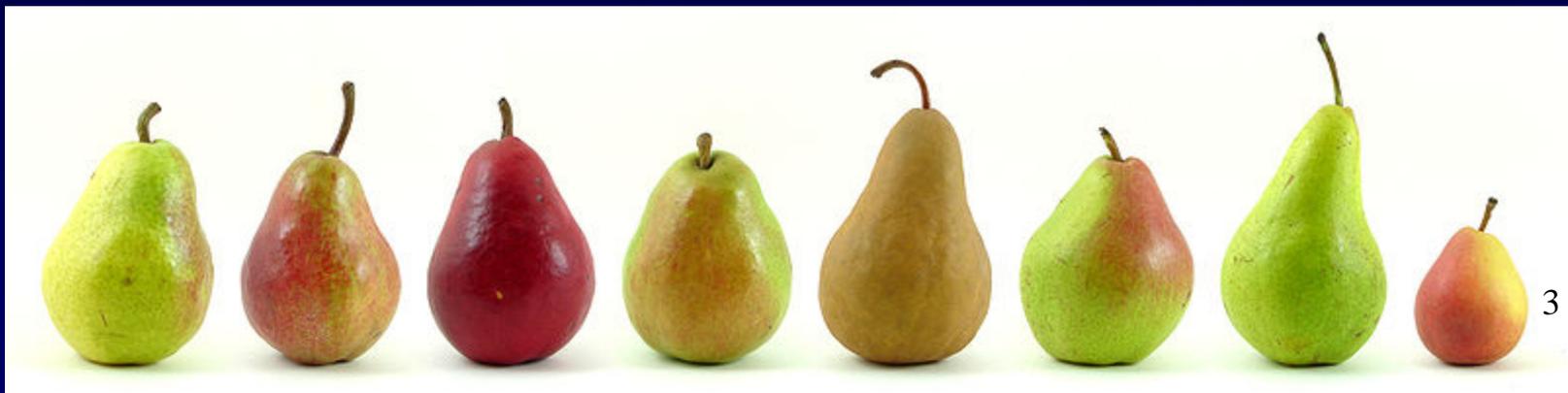
Beurré Hardy
Doyenne du Comice
Generale Leclerc
Passe Crassane

Intermediate

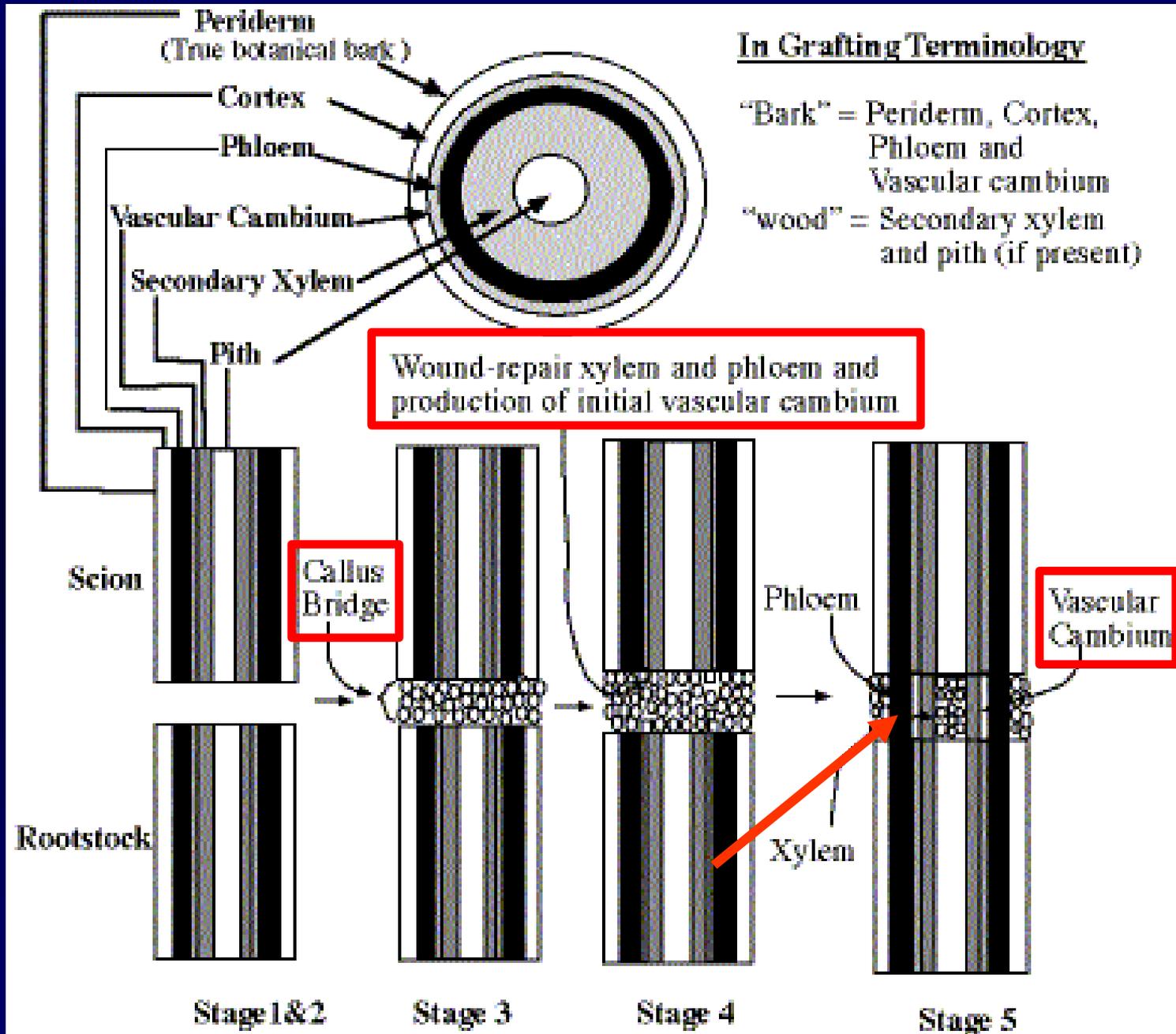
Abbé Fetel
Conference
Bartlett (William)
Packham's Triumph
Max Red Bartlett

Incompatible

Coscia
Clapp's Favorite
Dr. Guyot
Beurré Bosc_(Kaiser)



Graft histogenesis



- ◆ Initial phases of graft are the same in compatible or incompatible combinations, because they are a reaction to the wound.
- ◆ A good vascular tissue formation occurs only in the compatible combinations.

Graft Incompatibility

Scaramuzzi (1955): inability of some grafted plants to function as a single and unique plant.

Moore and Walker (1981): physiological incompatibility between tissues of the two members that leads to the failure of the graft.

Feucht (1988): phenomenon of premature tree senescence caused by physiological and biochemical dysfunctions.

Classification of graft-incompatibility:

- Translocated
- Delayed
- Localized

Translocated graft-incompatibility

- The **agent factor** is a substance transported from one graft member to the other as a **toxin**.
- The use of **interstock** graft can not overcome the incompatibility.
- Example: peach cv “Hale’s Early” grafted on “Myrobolan B” plum roots (weak union and distorted tissues).

Delayed graft-incompatibility

- After several years of growth, symptoms appear.
- This incompatibility is frequently due to the presence of disease, i.e. virus, phytoplasma, introduced by grafting.
- Incompatibility becomes evident when some mechanical stresses occur.

Quick decline



Abbé Fétel/ BA29 -Phytoplasma symptoms



Localized graft-incompatibility

- Physical contact between the two bionts are required.
- Other **specific symptoms** are necrosis vascular discontinuity, starch accumulation, break of the graft-union.
- Use of mutually compatible **interstock** overcomes the problem.
- An example: Beurré Bosc (Kaiser) grafted on quince → Beurré Hardy as interstock.

EXAMPLE OF PEAR-QUINCE

GRAFT-INCOMPATIBILITY

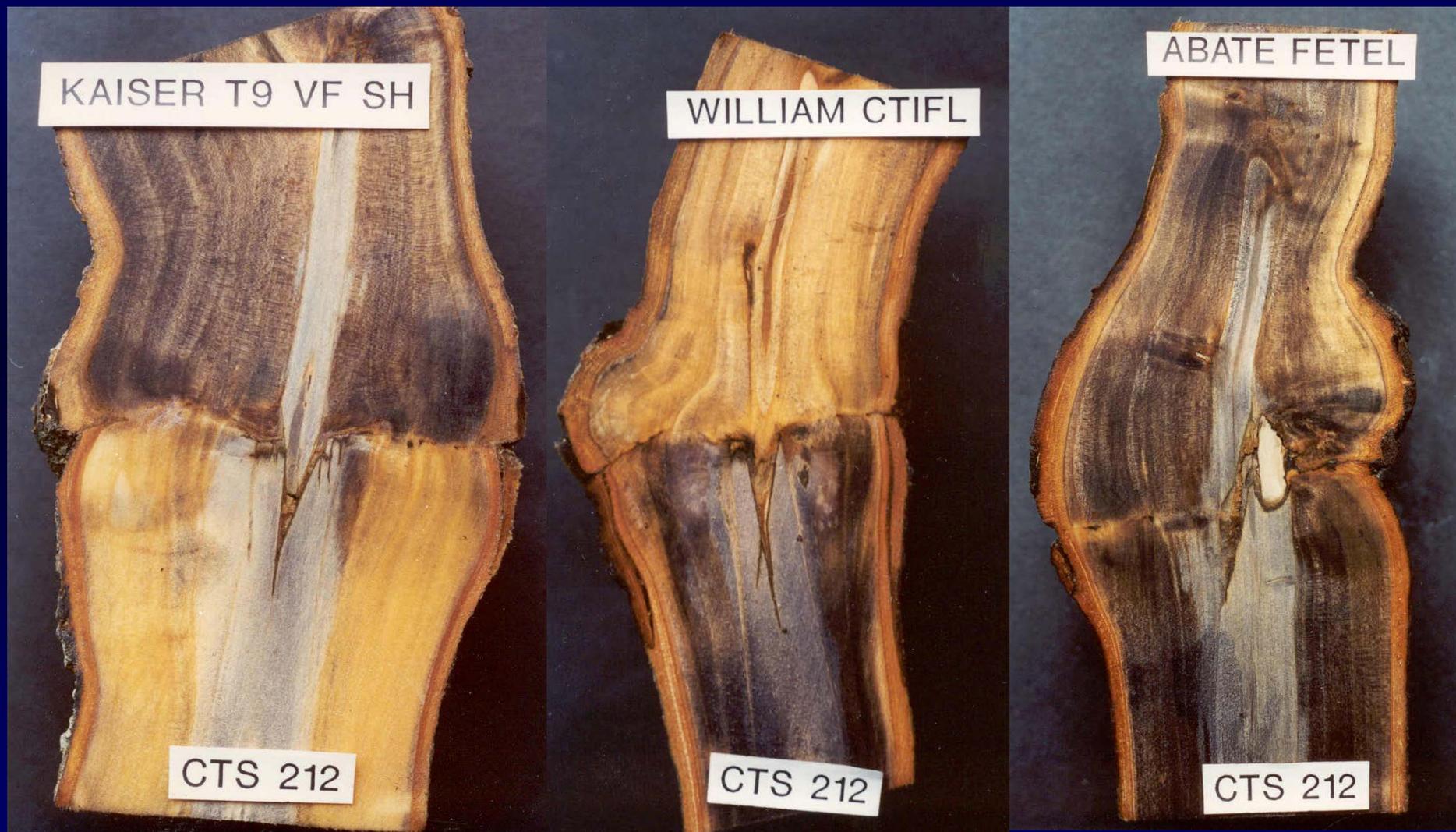
Bartlett

Necrosis
enlargement

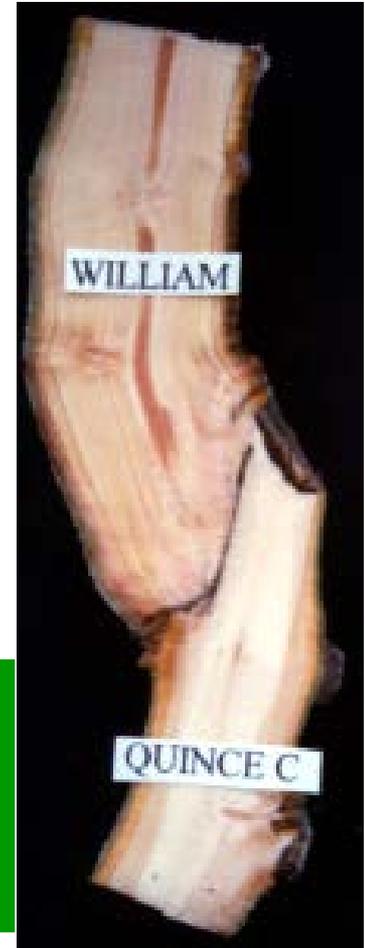
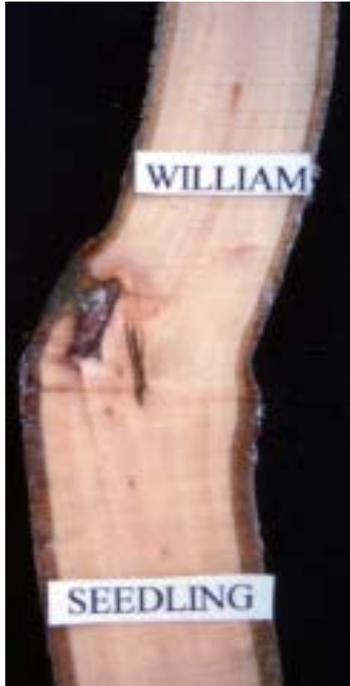
Quince C



Starch accumulation



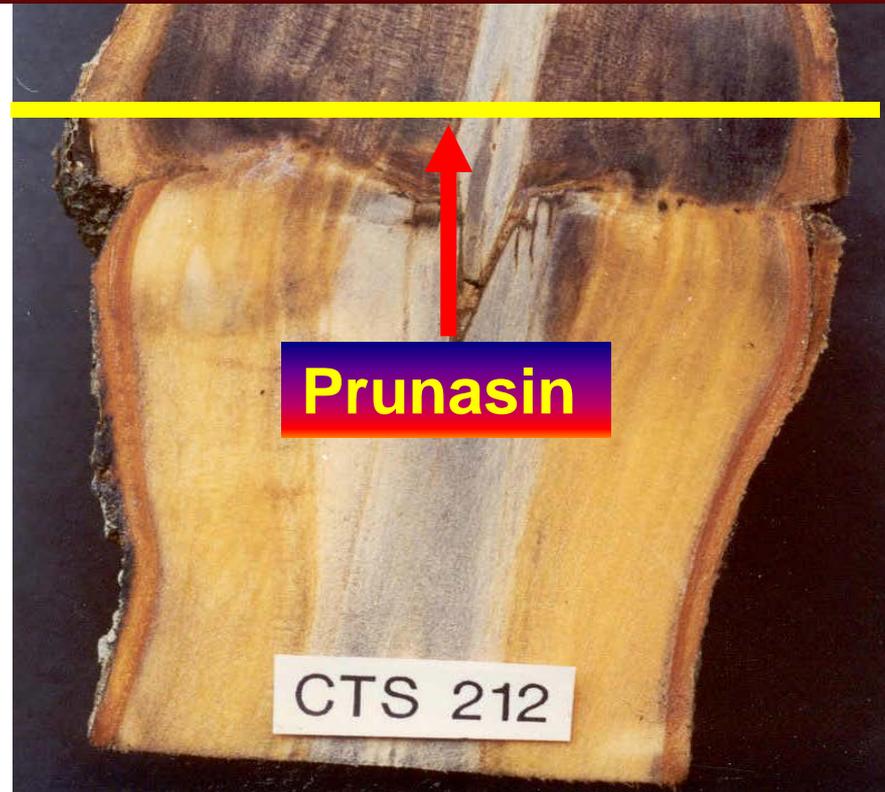
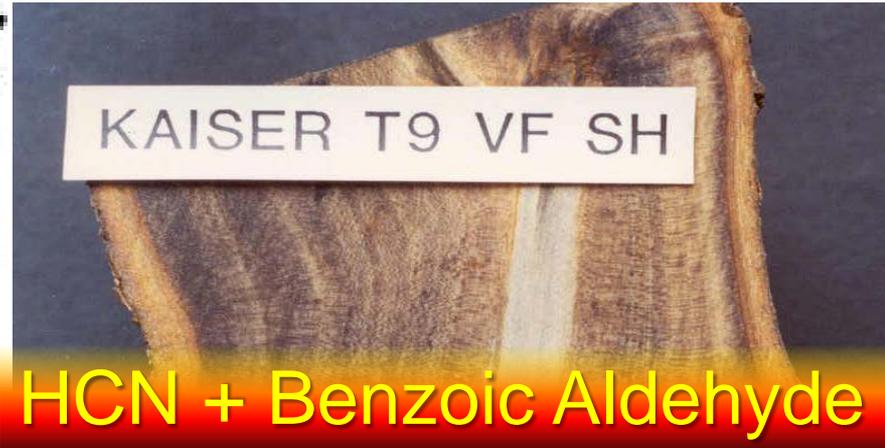
Graft-incompatibility



Gur's model of graft-incompatibility between pear and quince



Pear



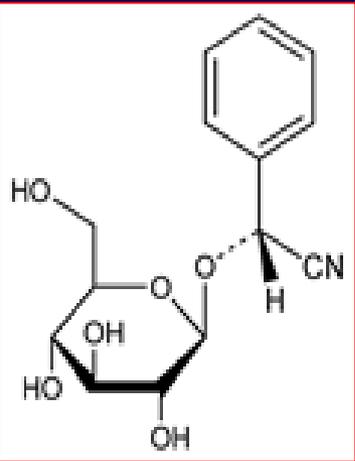
Quince



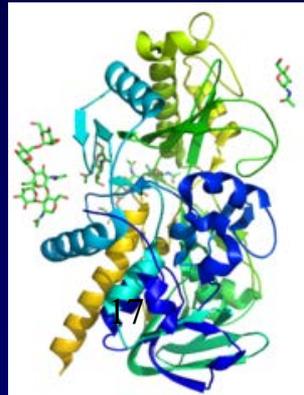
β -glucosidase
Mandelonitrile
lyase

Gur et al., 1968

In enzymology, a **prunasin beta-glucosidase** (EC 3.2.1.118) is an enzyme that catalyzes the chemical reaction:



The **mandelonitrile lyase** (EC 4.1.2.10) is an enzyme that catalyzes the chemical reaction:

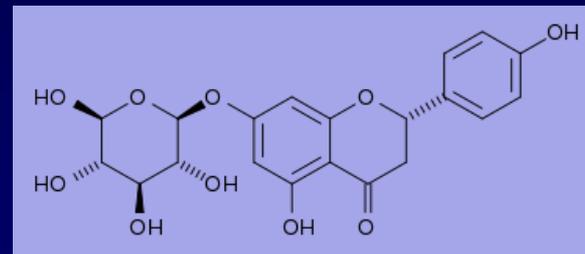


Objections to the Gur's model:

- Cyanogenic glycosides are accumulated and compartmentalised in cell vacuoles and are used as nitrogen reserve.
- There was no evidence of prunasin transportation **from quince to pear.**
- **There was no presence of prunasin in callus.**
- Cyanogenic glycosides are not present in pear.
- β -glucosidase and Mandelonitrile lyase are present also in the quince (detox system?)

Phenol involvement in Graft Incompatibility

Treutter and Feucht (1988) found an accumulation of prunin (*flavanon*) above the graft union in incompatible combinations of *Prunus avium* and *Prunus cerasus*, probably due to histo-anatomical disorders (e.g. changed cellular differentiation).



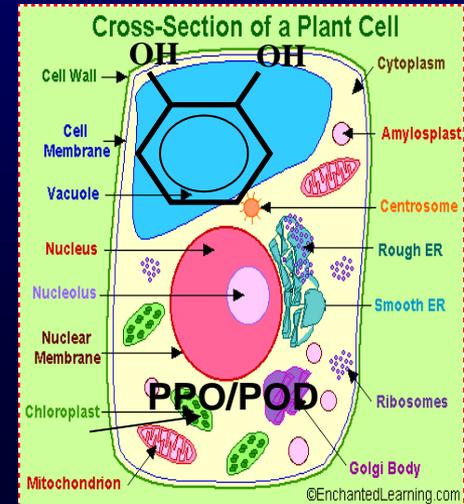
Phenol accumulation & graft-incompatibility

The different composition in polyphenols is known to **interfere with the plasmalemma** by changing the structure and the orientation of cell-wall microtubules and a corresponding modification of **wall cell permeability**.

(Errea, 1998)

Phenols in graft-incompatibility

- In graft incompatibility, the phenols may be exported from the vacuoles to the cytoplasmic matrix where they are oxidated by *PPO* and *POD* to quinones. The quinones polymerise and became toxic (Feucht and Treutter 1989).



- Moreover, they may form irreversible linkages with proteins or other macromolecules and precipitate with necrosis formation (Feucht and Treutter 1989).

Polyphenols found in the phloem of the different genotypes

Compound	Genotype
naringenina	William e Franco
+catechina	William, Franco e EMC
-epicatechina	William, Franco e EMC
quercitrina	William e Franco
isoquercitrina	EMC
procianidina B1	William
procianidina B2	EMC

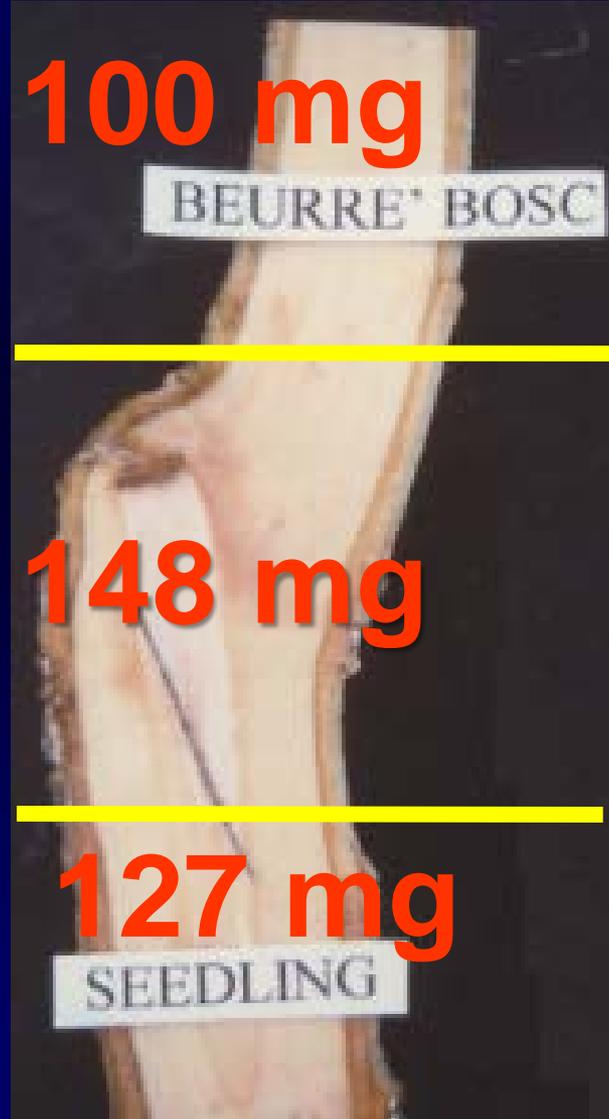
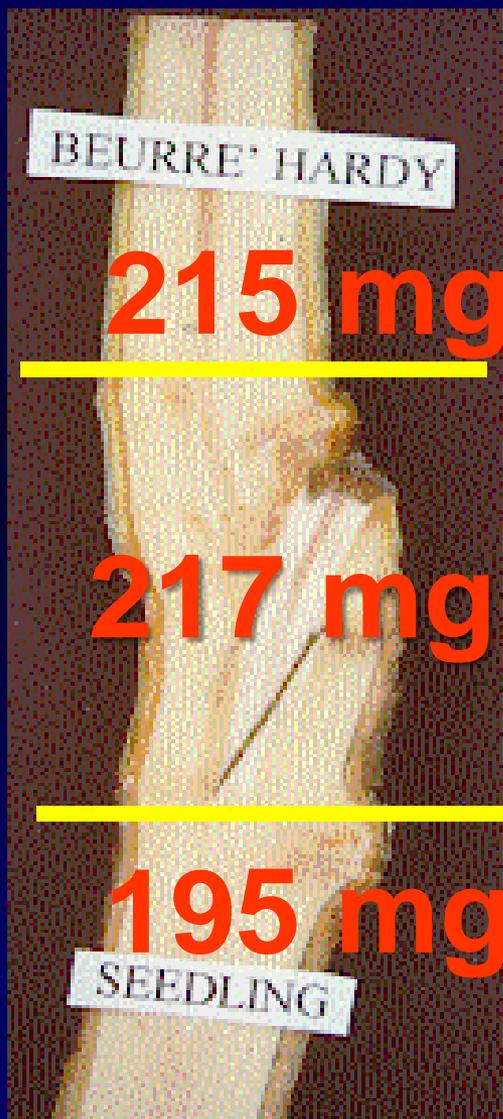
Polyphenol in the phloem

Combinations	Oligomeric fraction (%)
William autoradicato	0.78
Franco	0.95
Cotogno EMC	1.21
W/EMC 2 cm above	1.8
W/EMC Grafting point	2.9
W/EMC 2 cm below	1.4
W/Franco 2 cm above	0.9
W/Franco Grafting point	1.4
W/Franco 2 cm below	1.1

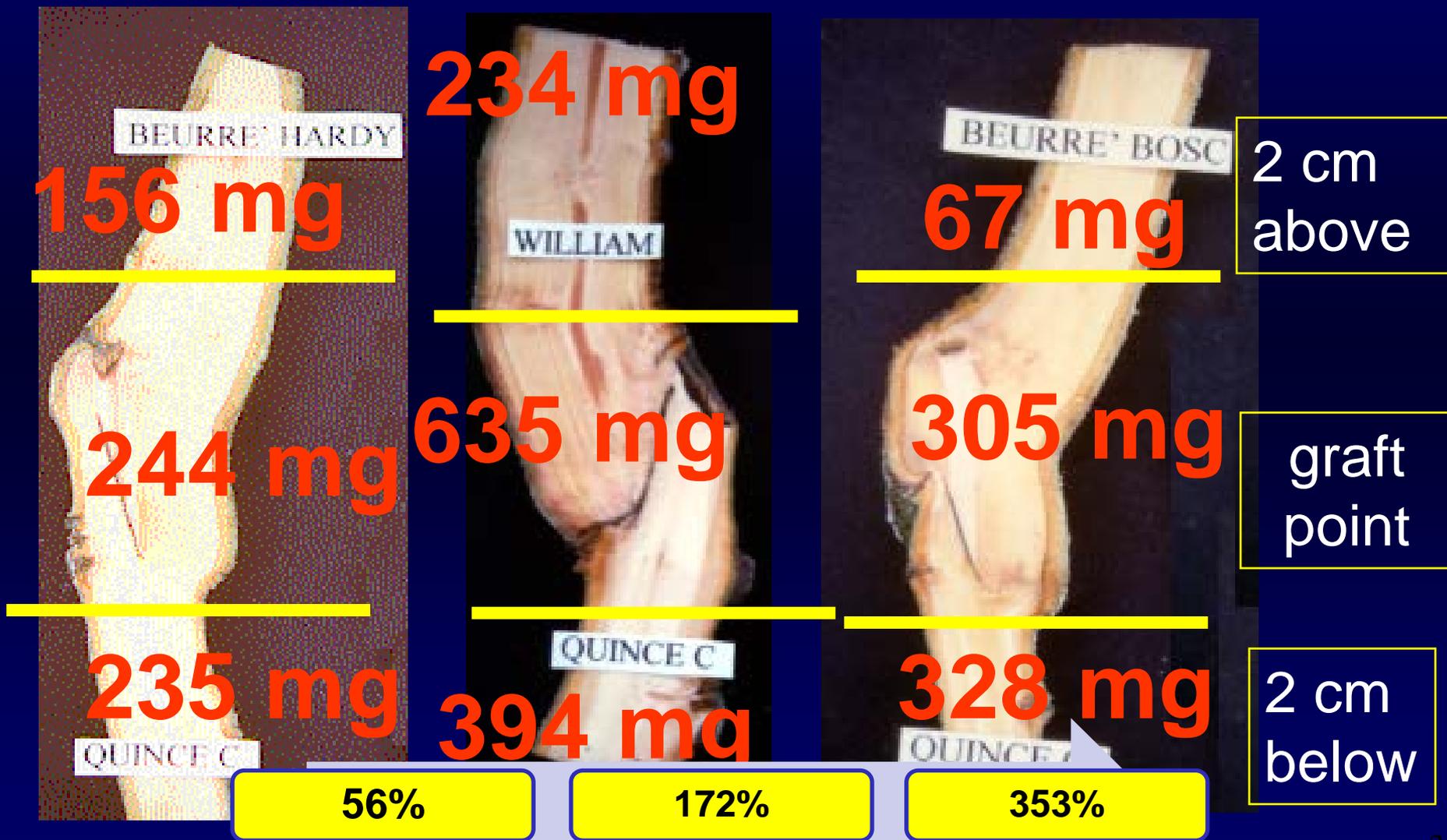
incompatible

compatible

PEAR-SEEDLING COMBINATION: (-)-EPICATECHIN (mg/100g FW)



PEAR-QUINCE COMBINATION: (-)-EPIGALLOCATHECHIN (mg/100g FW)



Phenols in pear-quince combinations

- Polyphenols accumulation at the graft point in incompatible combinations with quince was mainly due to (-)-epicatechin.
- This flavan-3-ol increased its concentration *at the graft union* compared with 2 cm above graft according to graft incompatibility levels with quince:
 - Beurré Hardy (+56%) ⇒ compatible
 - Bartlett (+172%) ⇒ intermediate
 - Beurré Bosc (+353%) ⇒ incompatible

Cell-cell recognition

Little information is available about cell-cell recognition, and **none** of the several theories posited to date **has established** the **existence of a recognition protein** between the cells of the two genotypes in the graft union tissues.

Yeoman and Brown (1976)

Proposed a model based on the formation of a cell-cell recognition **protein complex** for the scion-stock combination: its absence is supposed to demonstrate an incompatible graft.

Moore (1981-1984)

Using “in vitro” callus models, he showed that contact is not necessary for the onset of graft incompatibility.

How to study graft-incompatibility?

- *In vivo (polyphenols role)*
- *In vitro (new models to simplify the system and reduce interactions with the environment)*

Models for studying biochemical and physiological aspects of graft-incompatibility



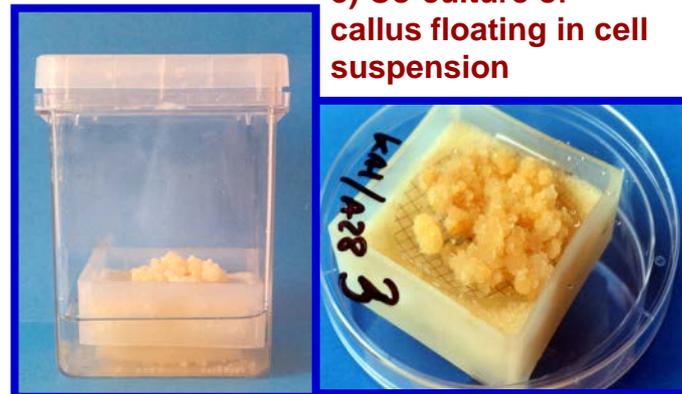
a) Micrografting *in vitro*



c) Co-culture of callus in the same Petri dishes



b) Graft of *in vitro* shoot on acclimating plants



e) Co-culture of callus floating in cell suspension



d) Co-culture of cell suspensions

A) *In vitro* Micrografts



BUTIRRA HARDY

BEURRE HARDY

KAISER

BEURRE BOSCH

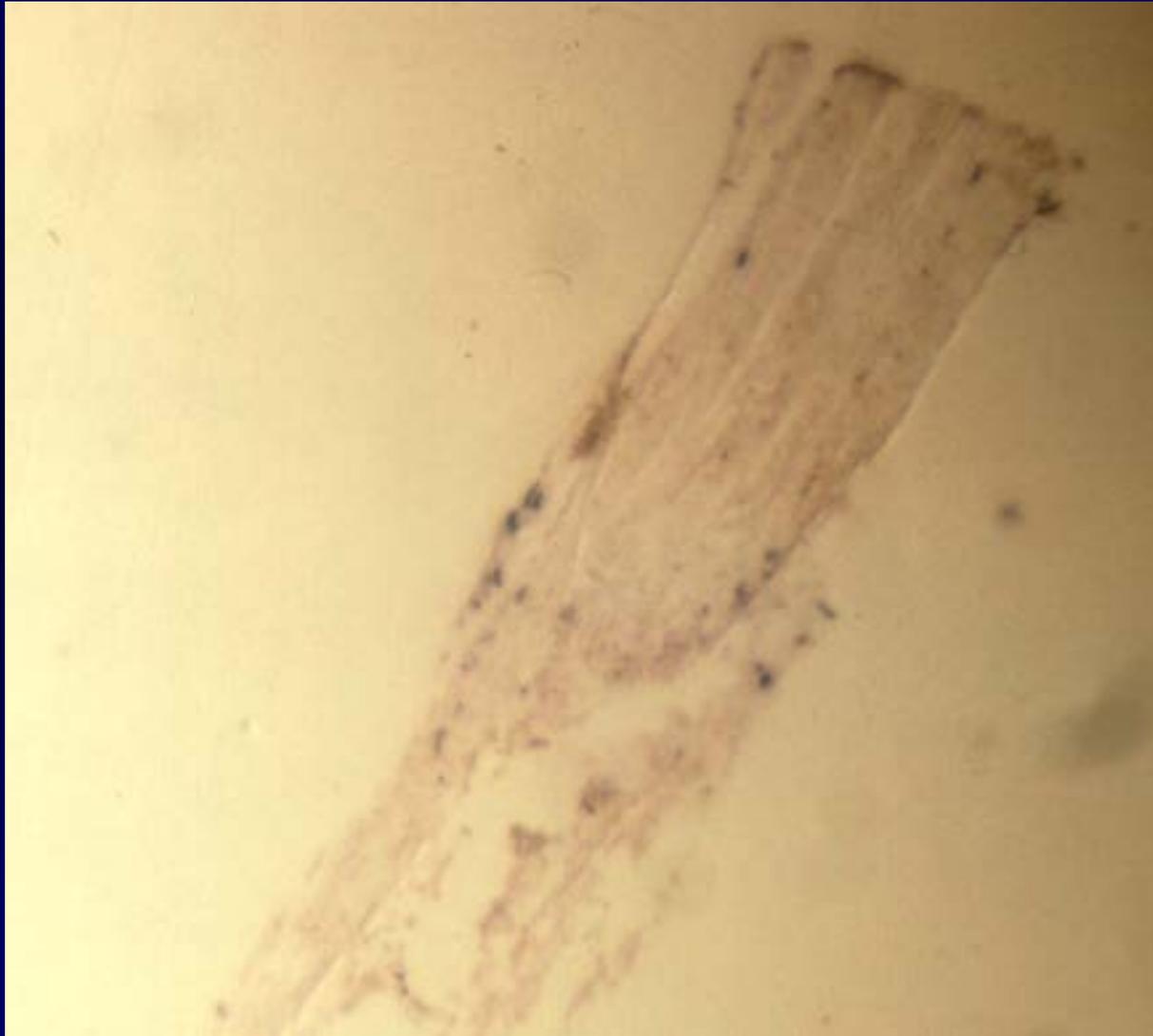
WILLIAM

BARTLETT



EM C

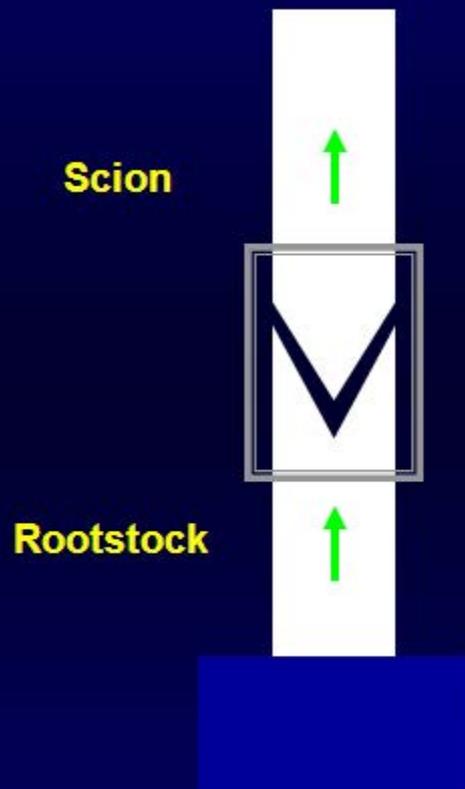
Tissue printing to determine presence of enzyme (beta glucosidase)



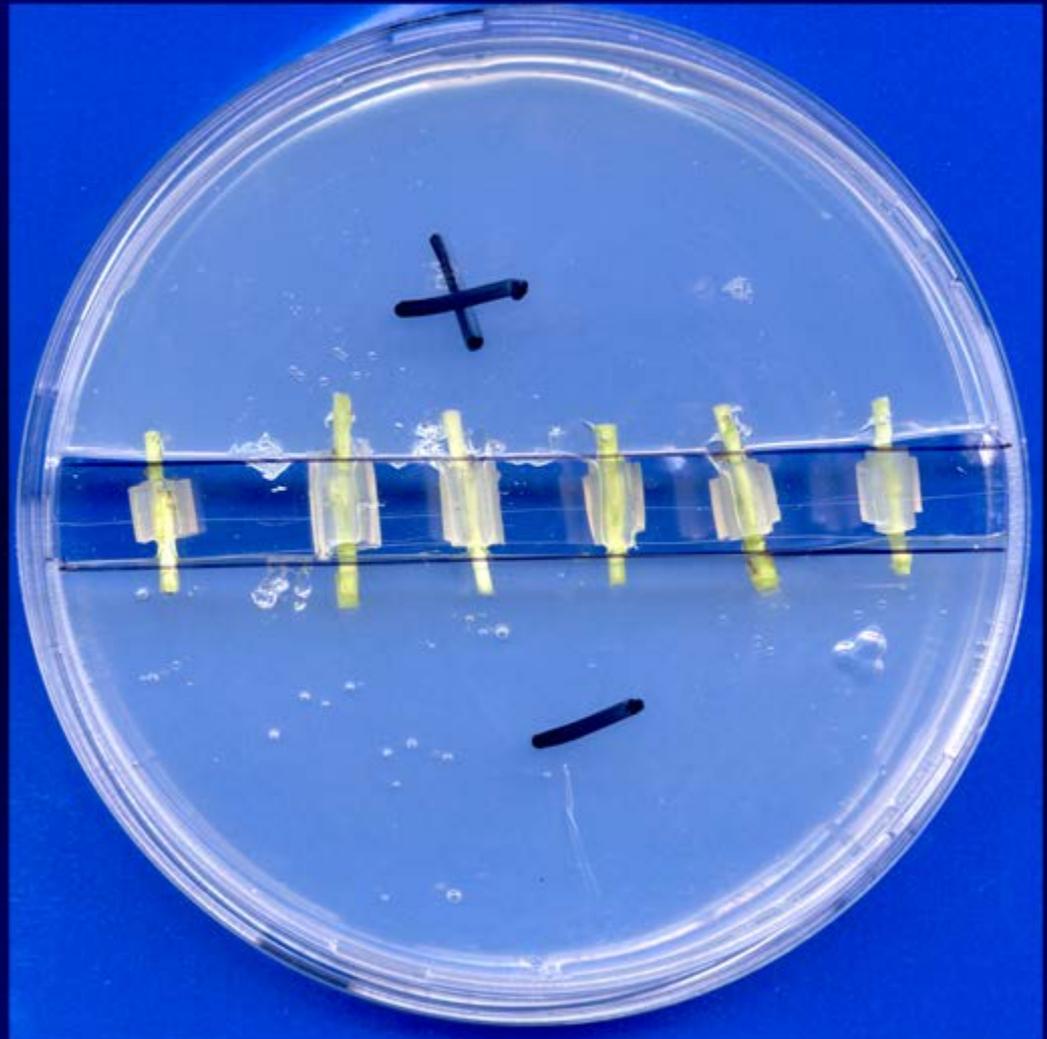
Tissue printing to determine presence of beta glucosidase



In vitro micrografts



Bosc/Bosc Bosc/MC
BH/BH BH/MC



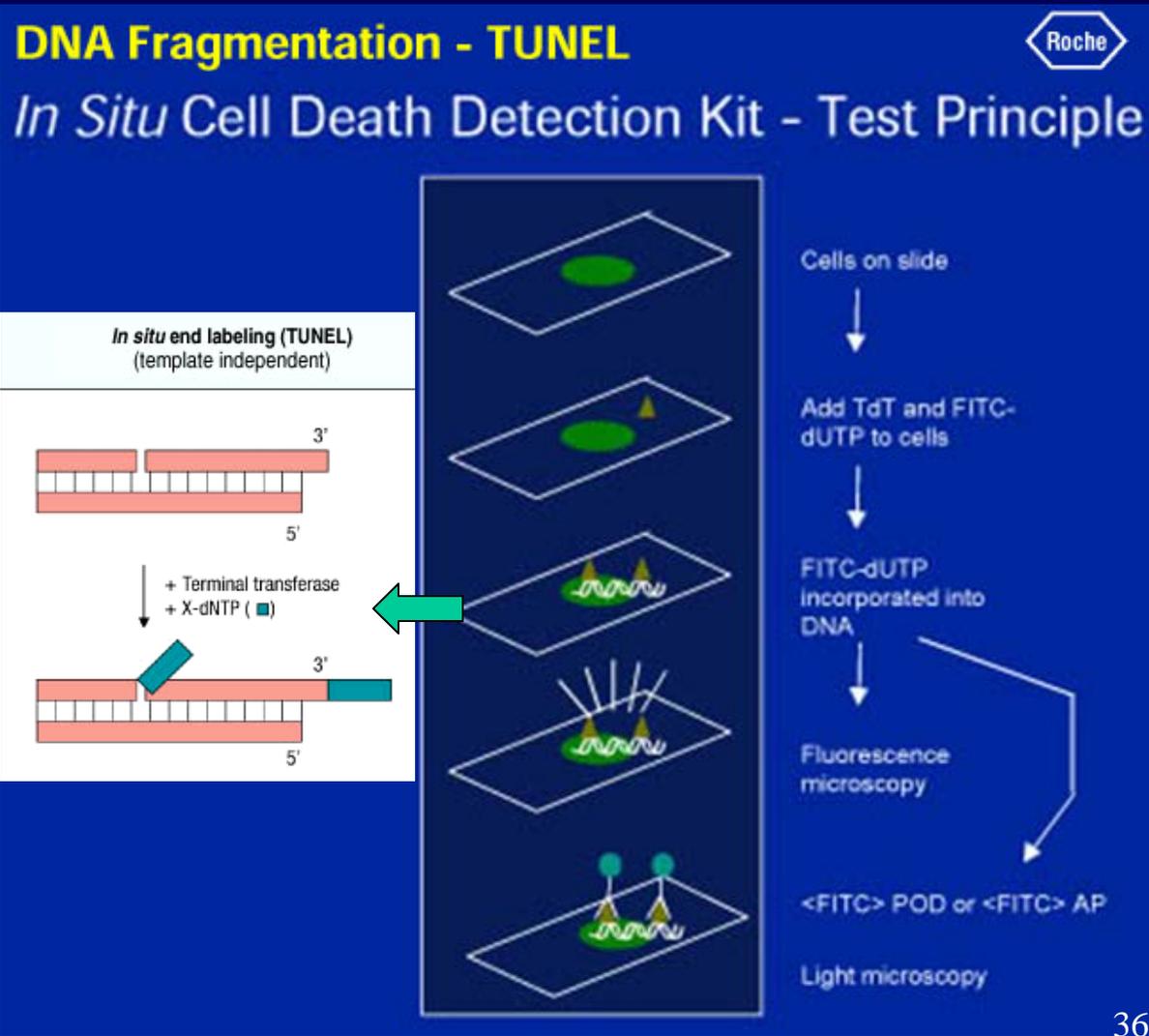
Espen et al., 2002 and 2005

The *In Situ Cell Death Detection Kit* [Roche Diagnostics, GmbH] has been designed as a precise, fast, and simple non radioactive technique to detect and quantify apoptotic cell death at the single-cell level in tissues; based on the **TUNEL method (TdT-mediated dUTP terminal nick-end labeling to detect PCD-characteristic DNA fragmentation)**

During apoptosis, DNase activity not only generates double-stranded, low-molecular-weight DNA fragments (mono- and oligonucleosomes), but also introduces strand breaks ("nicks") into the high-molecular-weight DNA.

These processes can be identified by labeling the free 3'-OH termini with terminal transferase (TdT), which attaches labeled nucleotides to all 3'-OH-ends (TUNEL reaction; TdT-mediated dUTP nick end labeling).

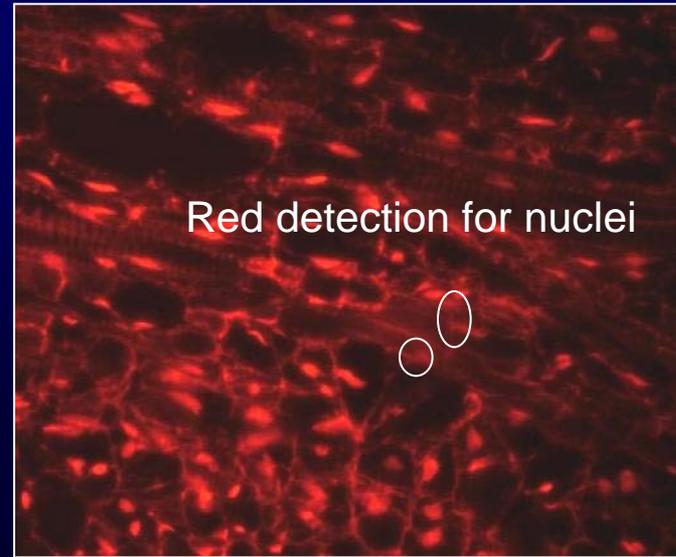
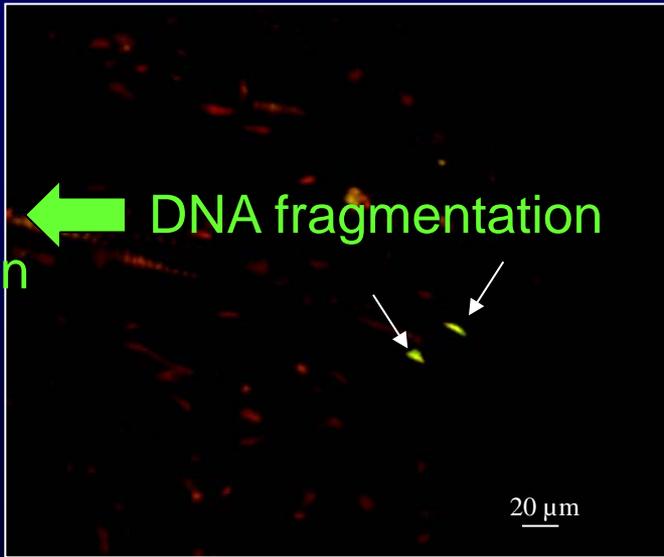
Labeling with fluorescein may also be followed by immunohistochemical detection using anti-fluorescein-specific antibodies that are conjugated to POD or AP.



***In situ* TUNEL assay:**

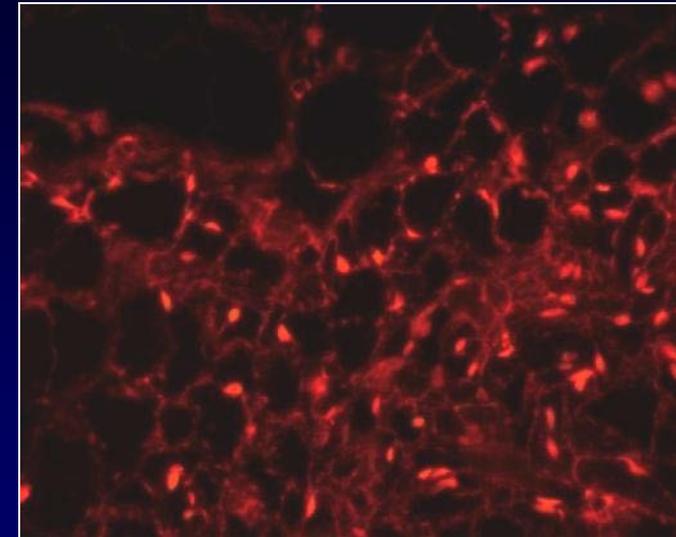
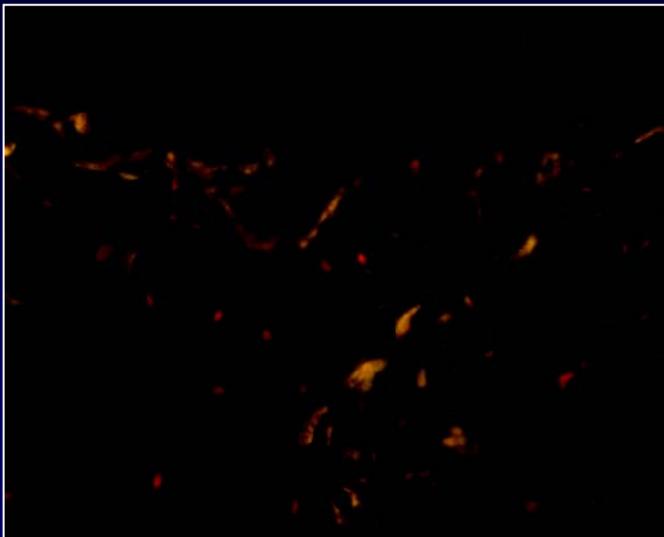
TdT-mediated dUTP terminal nick-end labeling to detect PCD-characteristic DNA fragmentation

TE
differentiation



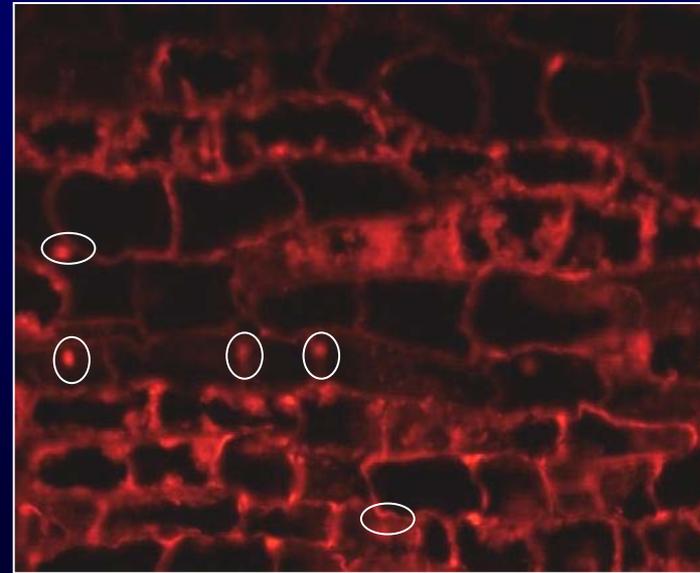
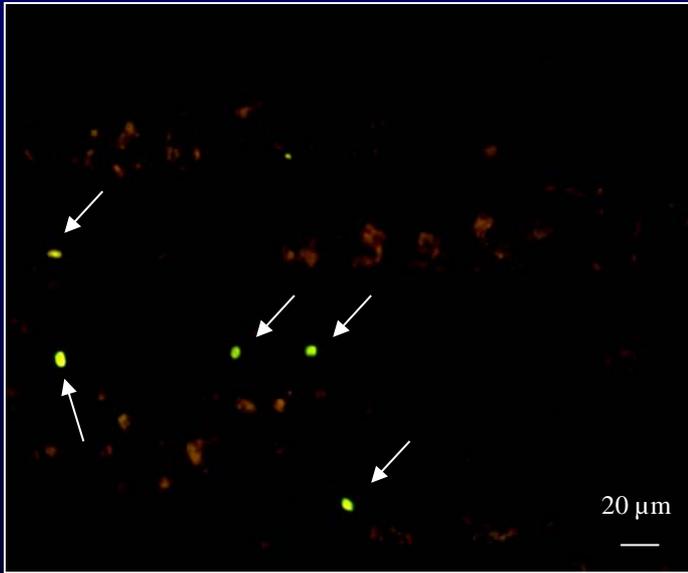
10 DAG

Bosc/Bosc: compatible



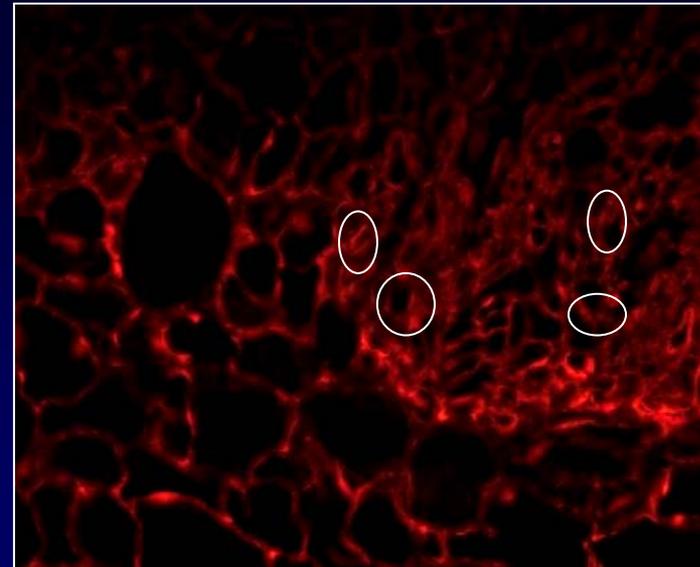
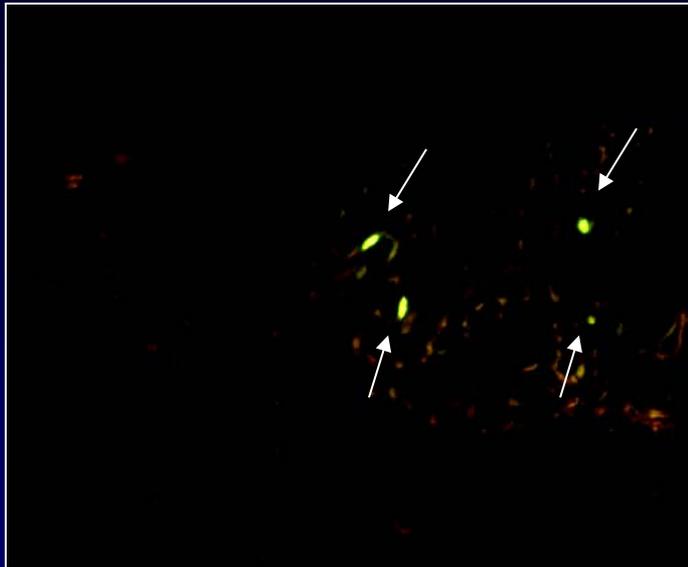
Bosc/MC: incompatible

In situ TUNEL assay



10 DAG

BH/BH: compatible



BH/MC: compatible

b) Graft of in vitro shoot on acclimating plants

Grafting phases



A) Rootstock preparation.



B) In vitro shoot.



C) Grafted plant

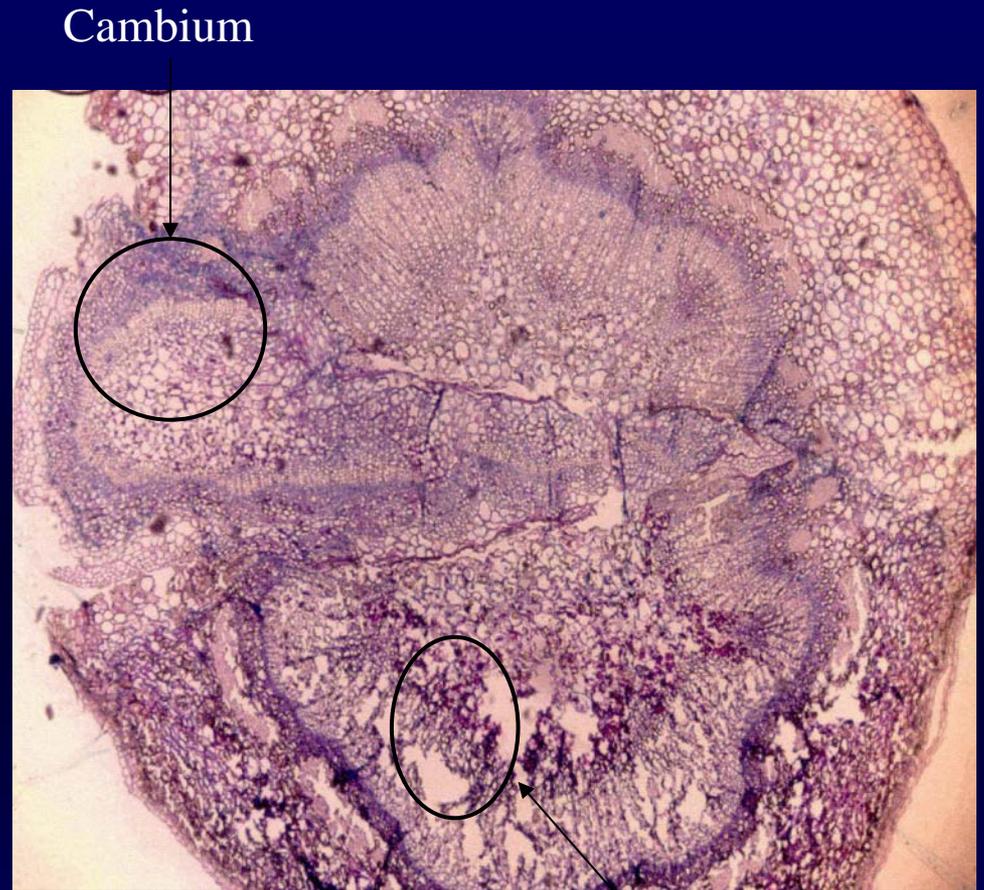


D) Protection to avoid shoot drought.



K/OHF after 30 days

**Merged
cambium**



BH/OHF after 30 days

Starch

Cambium

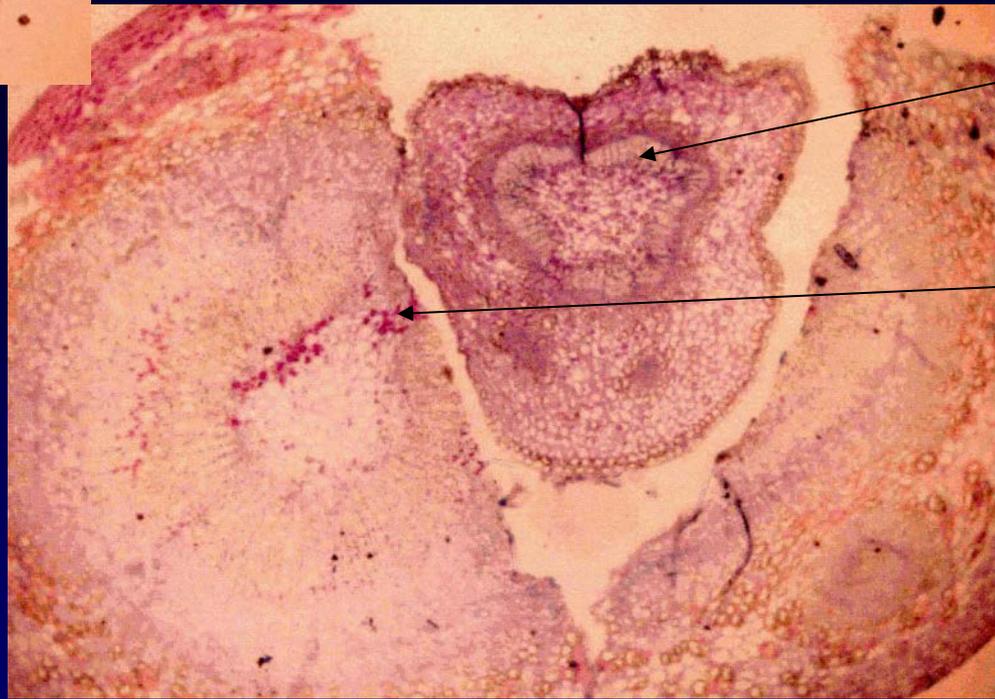


Rootstock

Callus

BH/MC after 30 days

Starch



Cambium

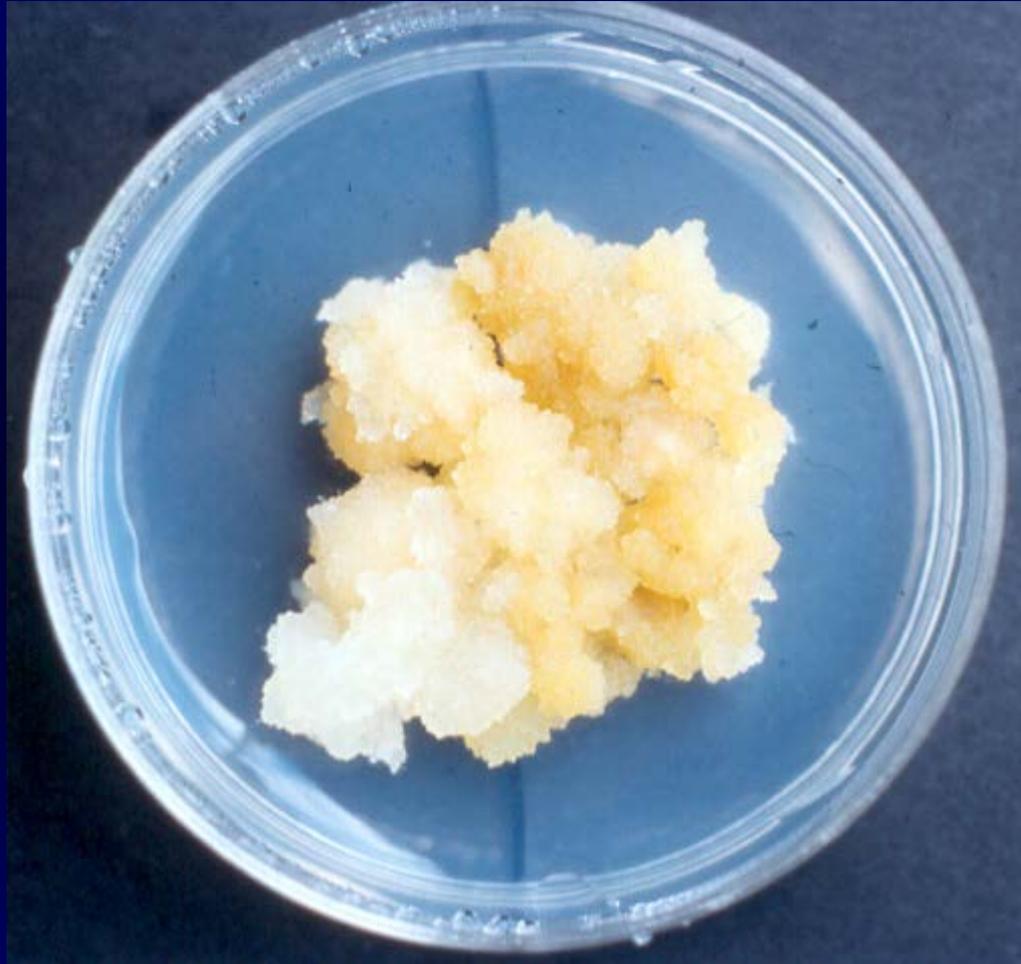
Starch

K/MC after 30 days

c) Co-culture of callus in the same Petri dishes

Example 1

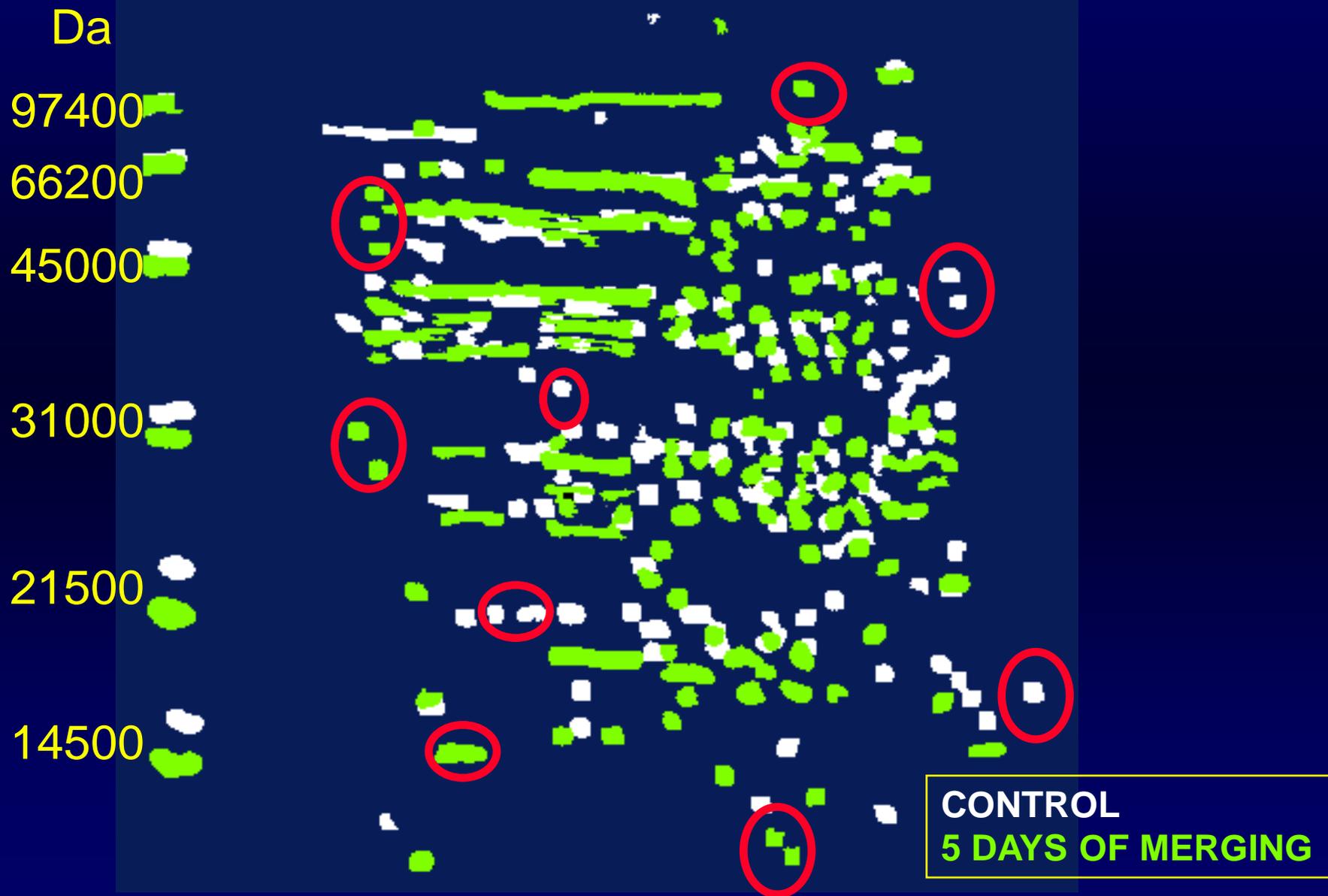
Bartlett



**Quince
BA29**

Musacchi et al., 1996

PROTEINS COMPARISON OF BEURRE' HARDY AND QUINCE C CALLUS CONTROL (NEVER MERGED) AND AFTER 5 DAYS OF MERGING



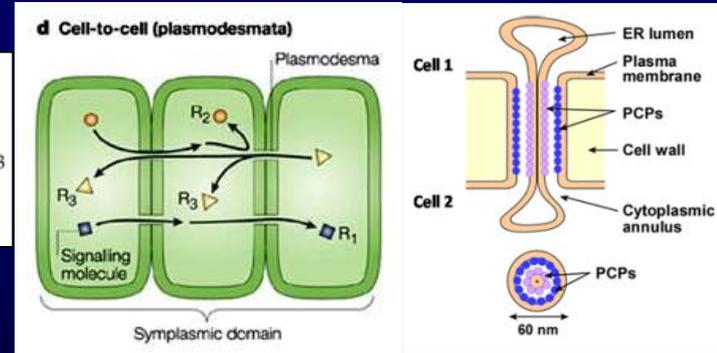
c) Co-culture of callus in the same Petri dishes

Example 2

Cell-to-cell transport through plasmodesmata in tree callus cultures

ANA PINA,^{1,2} PILAR ERREA,¹ ALEXANDER SCHULZ² and HELLE J. MARTENS^{2,3}

Tree Physiology 29, 809–818 2009



Starting point:

- for a successful grafting, the establishment of symplastic contacts in graft interface facilitates compounds transfer between 2 bionts.
- Plasmodesmata mediate the cell-to-cell communication route in the plant kingdom

Hypothesis:

- Localized incompatibility (in some *Prunus* grafts) could be related to insufficient plasmodesmata coupling at an early stage of development within one of the partners. Idea to be verified through bioimaging methods.

Material:

MO callus= apricot cv. Monique

MN callus= plum rootstock Marianna 2624

MO/MO and **MN/MN** → compatible homografts

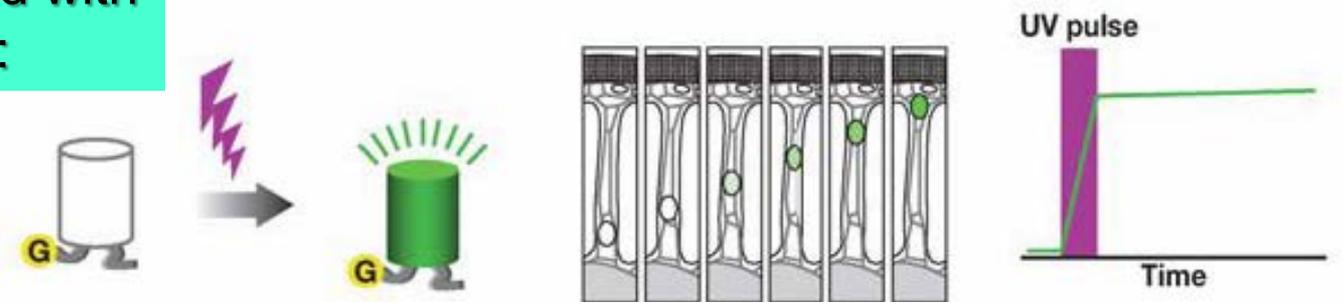
MO/MN → incompatible heterograft



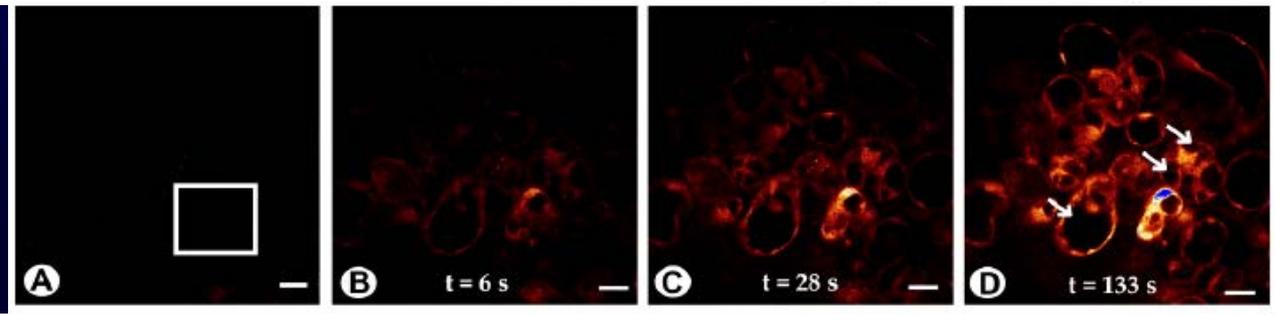
Callus heterograft

Techniques combined with confocal microscopy:

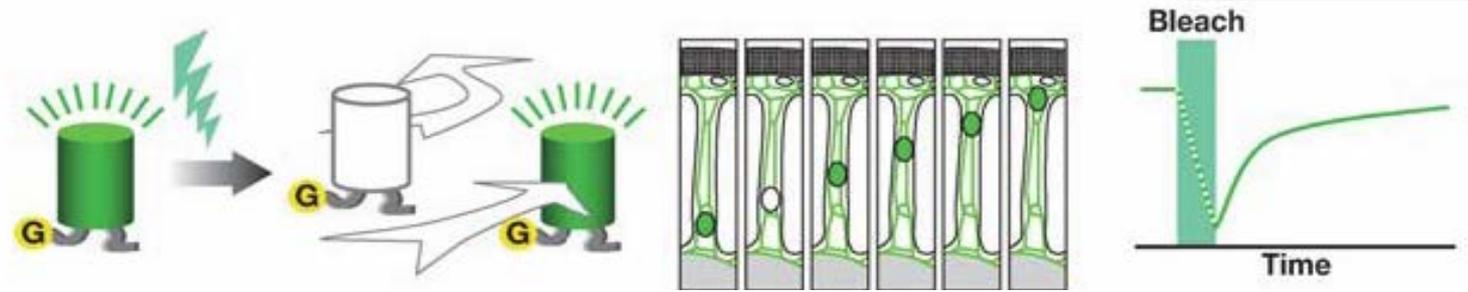
Photoactivation (PAF)



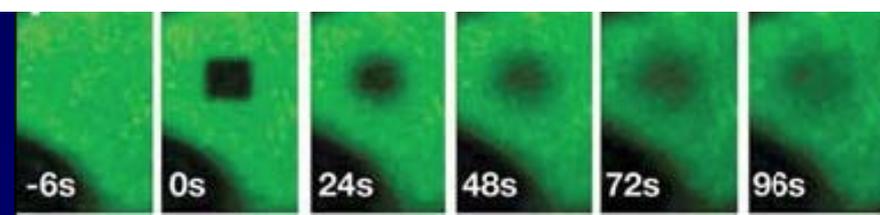
To track the diffusion of released fluorescein via plasmodesmata

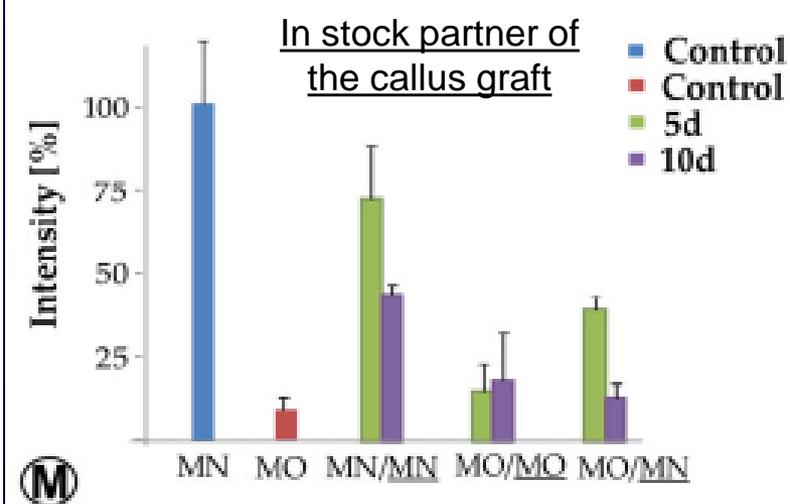
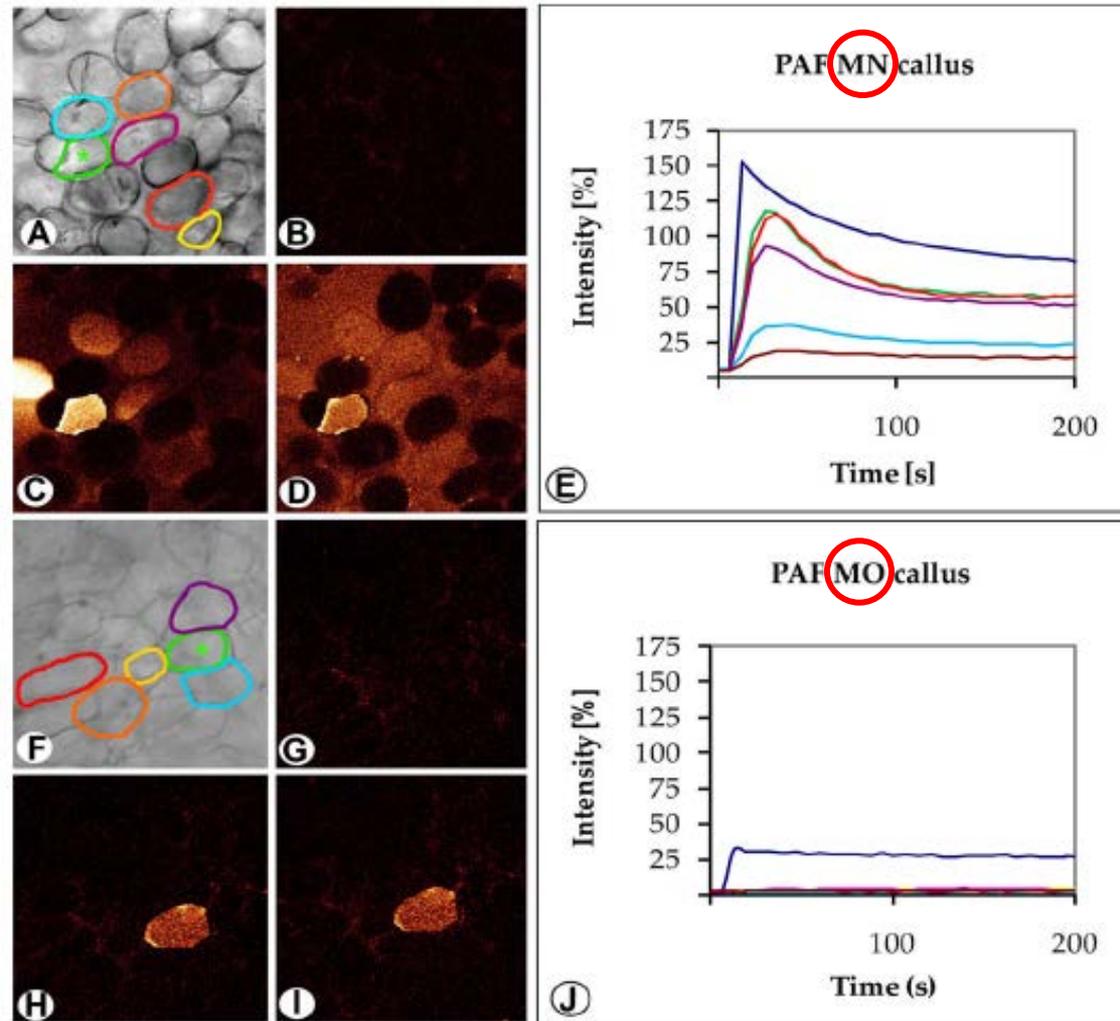


Fluorescence recovery after photobleaching (FRAP)



To quantitatively measure the movements of fluorescently tagged molecules or structure within live cells after photobleaching





FRAP quantitative experiments confirm the results of a **poor rate of symplastic communication in the apricot MO cv.** as found with PAF. This may affect the passages of nutrients and macromolecules at early stages of developmental between MO and MN in heterograft.

Novel control factor of connectivity that reaches the graft partner and alters its innate rate of communication.

d) co-culture of cell suspensions





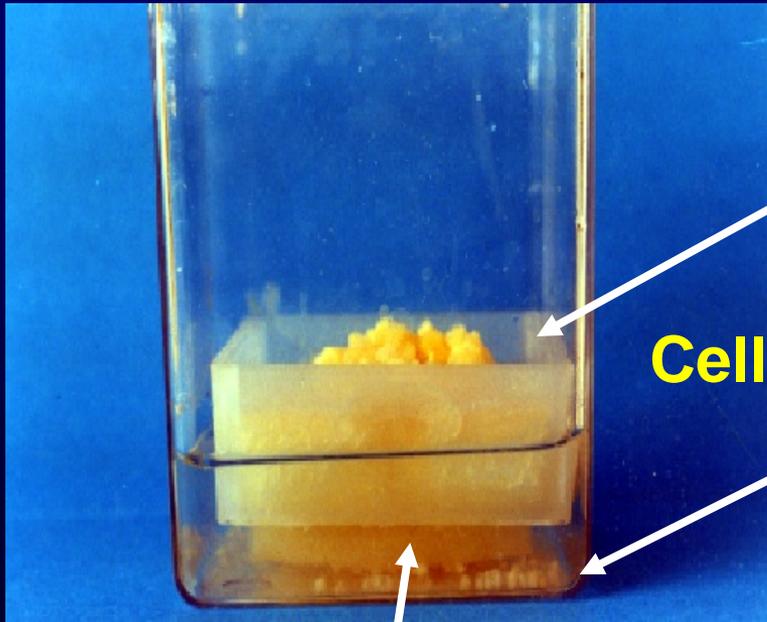
Dark, $\sim 22^{\circ}\text{C}$ for 14 days at 78 rpm

e) co-culture of callus floating in cell suspension

A “Magenta” and a raft with a low protein-absorption membrane (Durapore, 5 μm pore size, Millipore)

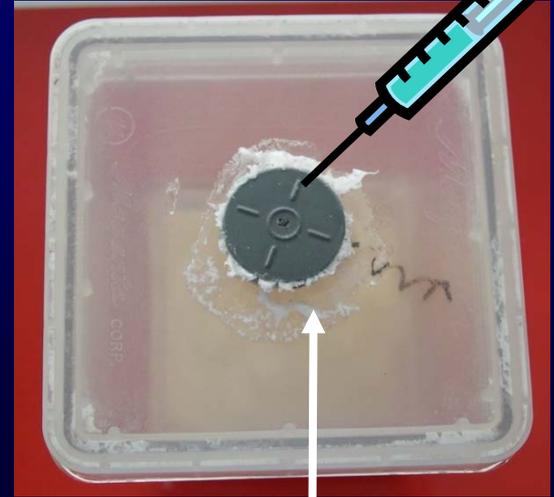
The graft union is simulated by placing the callus over the membrane and the cell-suspension cultures under it, so that the raft, kept in motion by a mechanical stirrer, floats.





Callus

Cells suspension



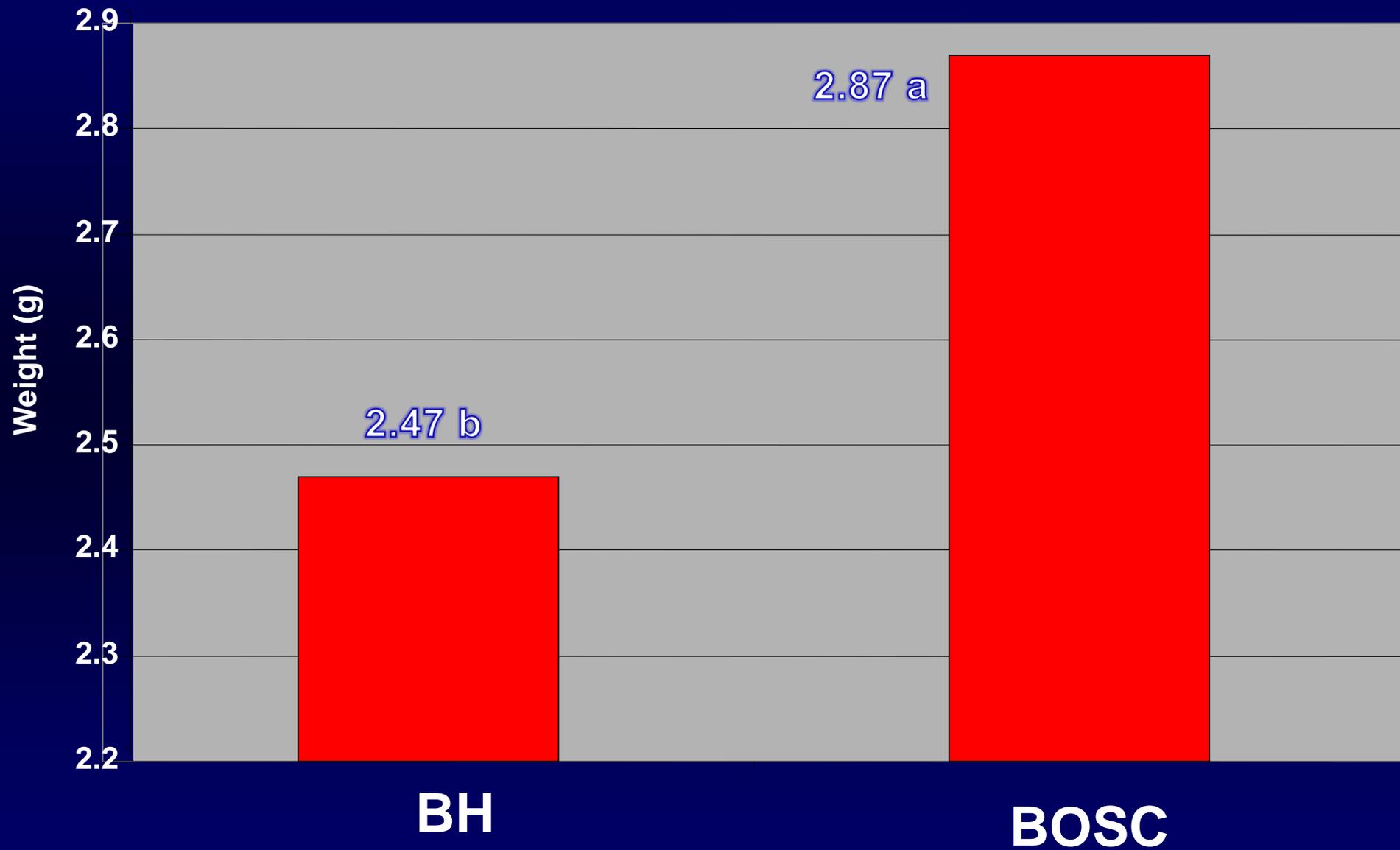
**Magenta lid with
rubber septum**



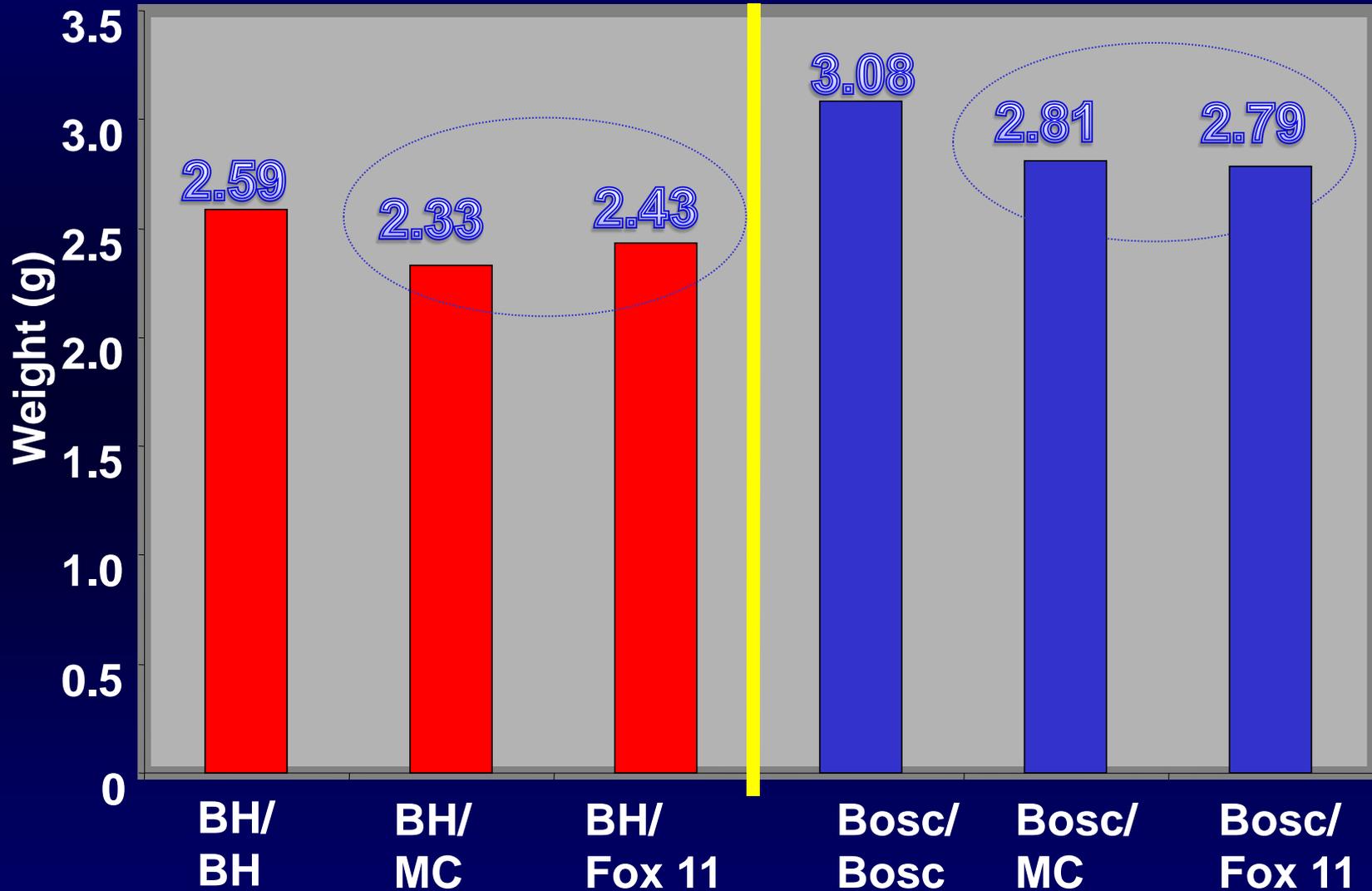
Sponge

Membrane

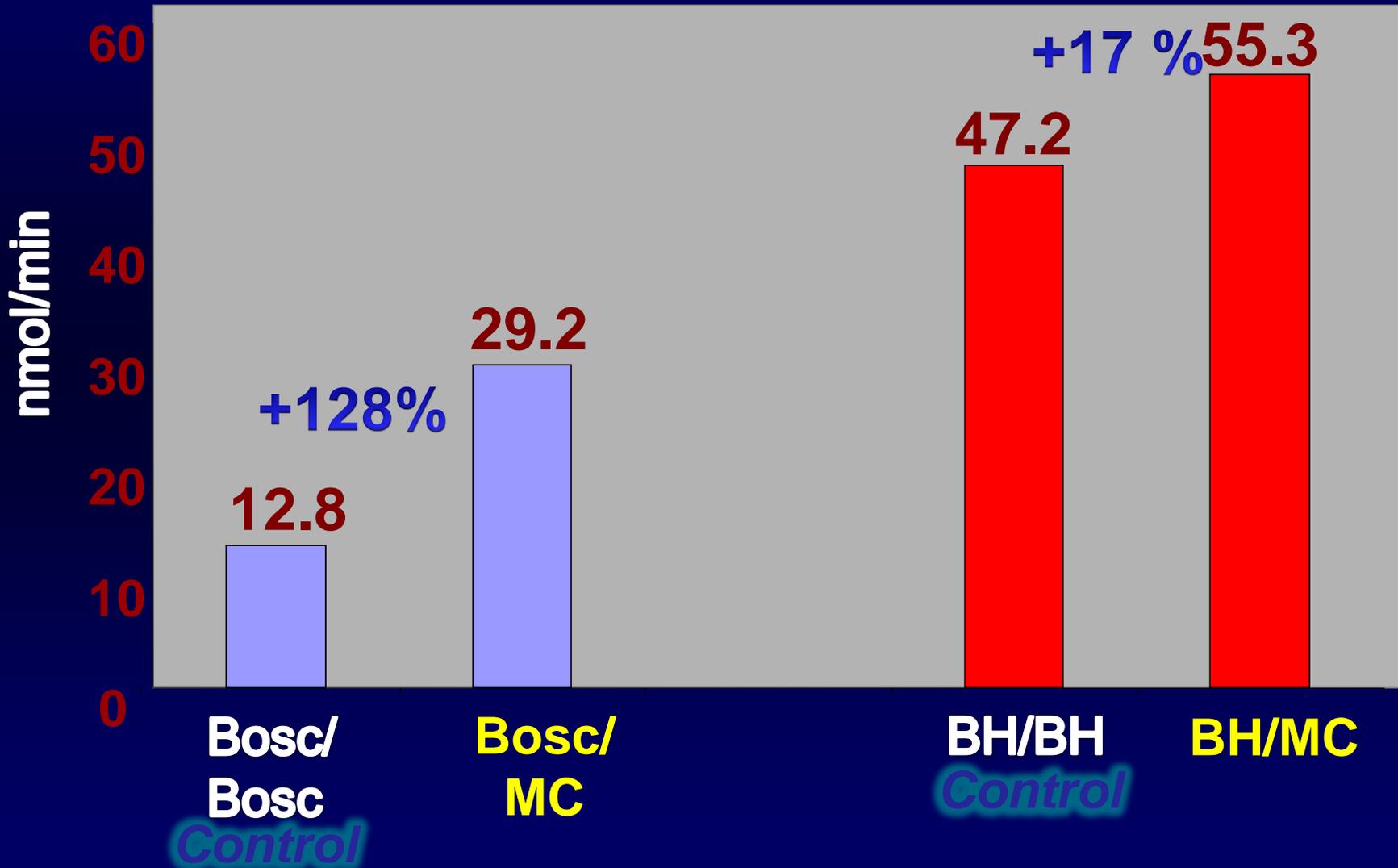
Cultivar callus growth



Callus growth after one week of co-culture

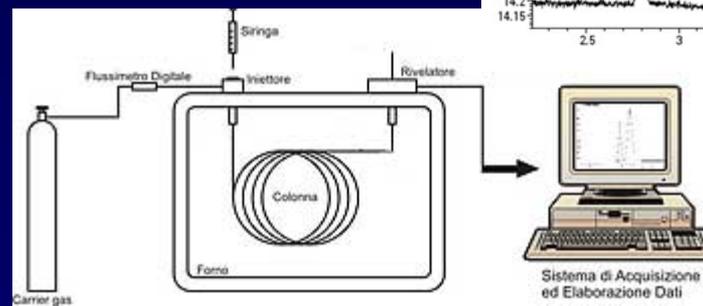
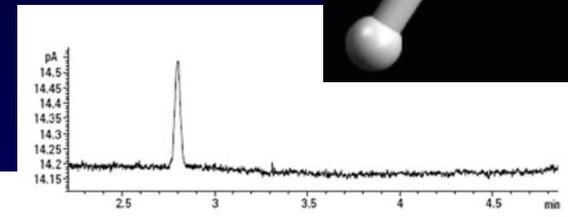
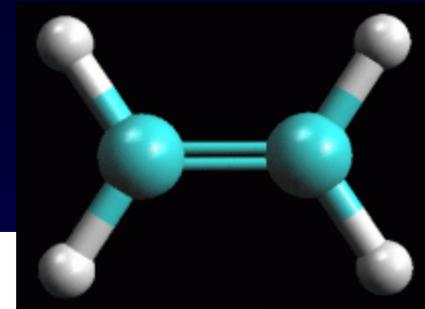


Respiration rate

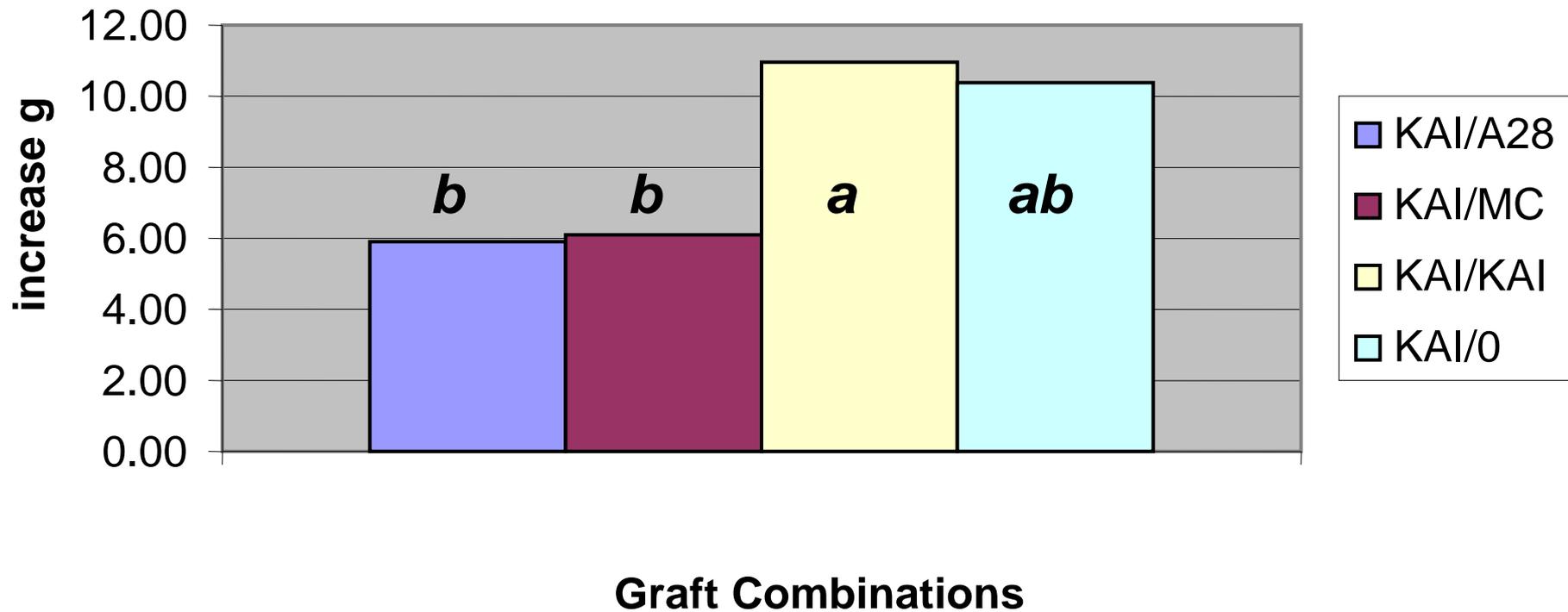


GAS ANALYSIS

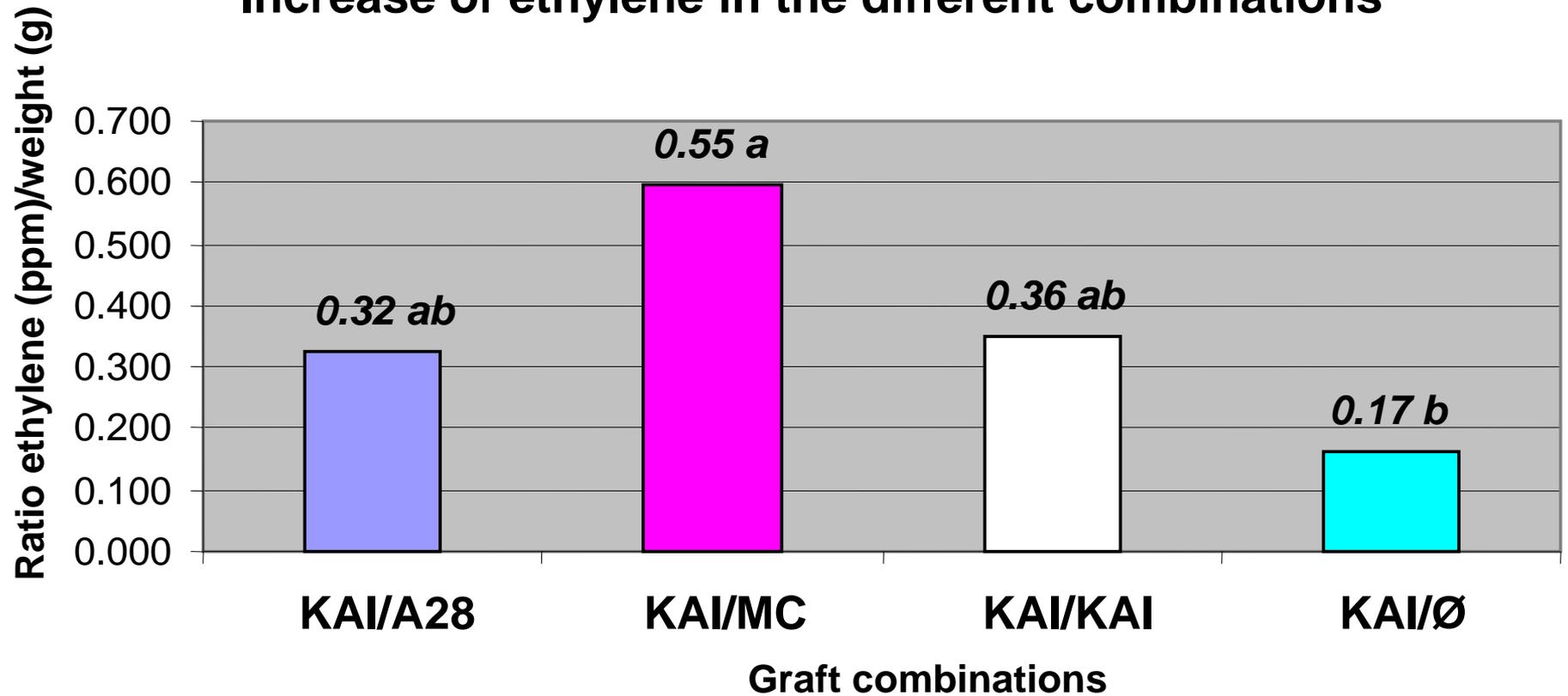
- CO₂
- Ethylene



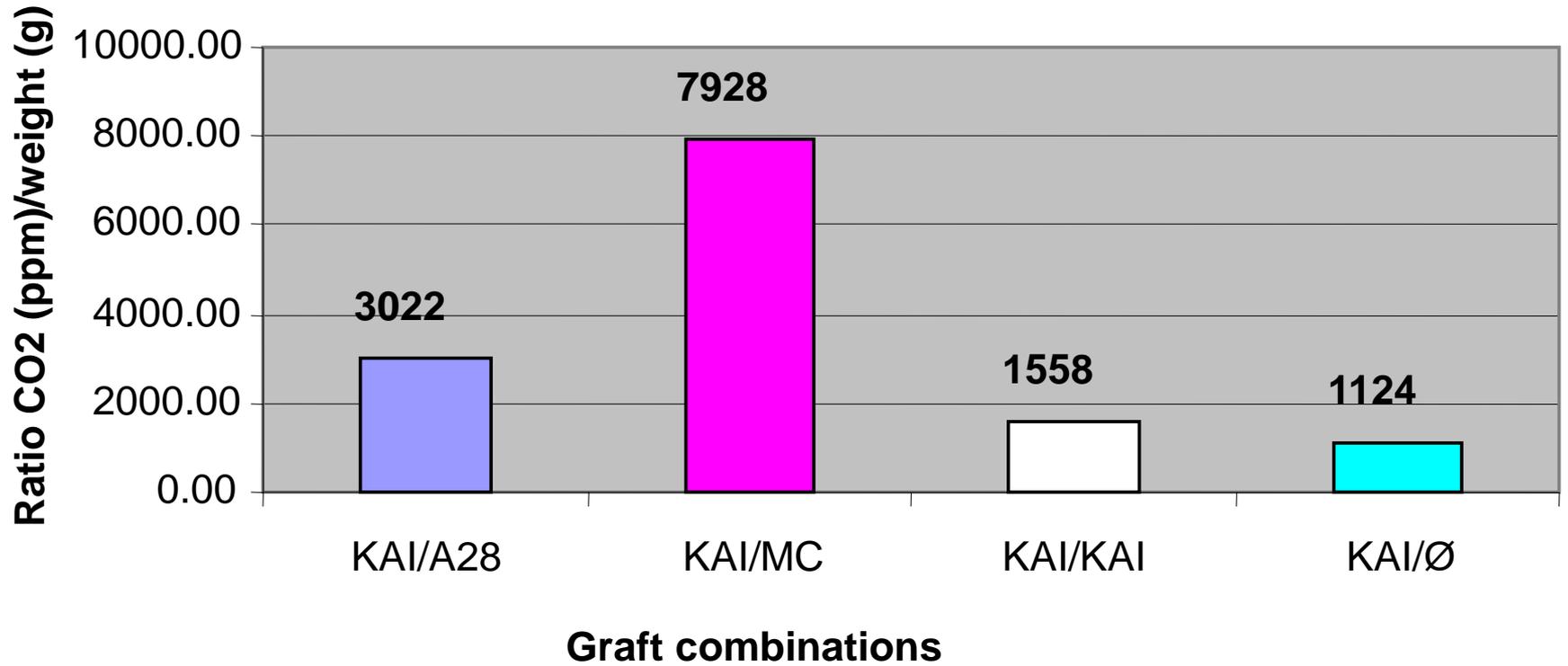
Bosc – increase of callus weight (g)



Increase of ethylene in the different combinations



CO₂ increase in different combinations

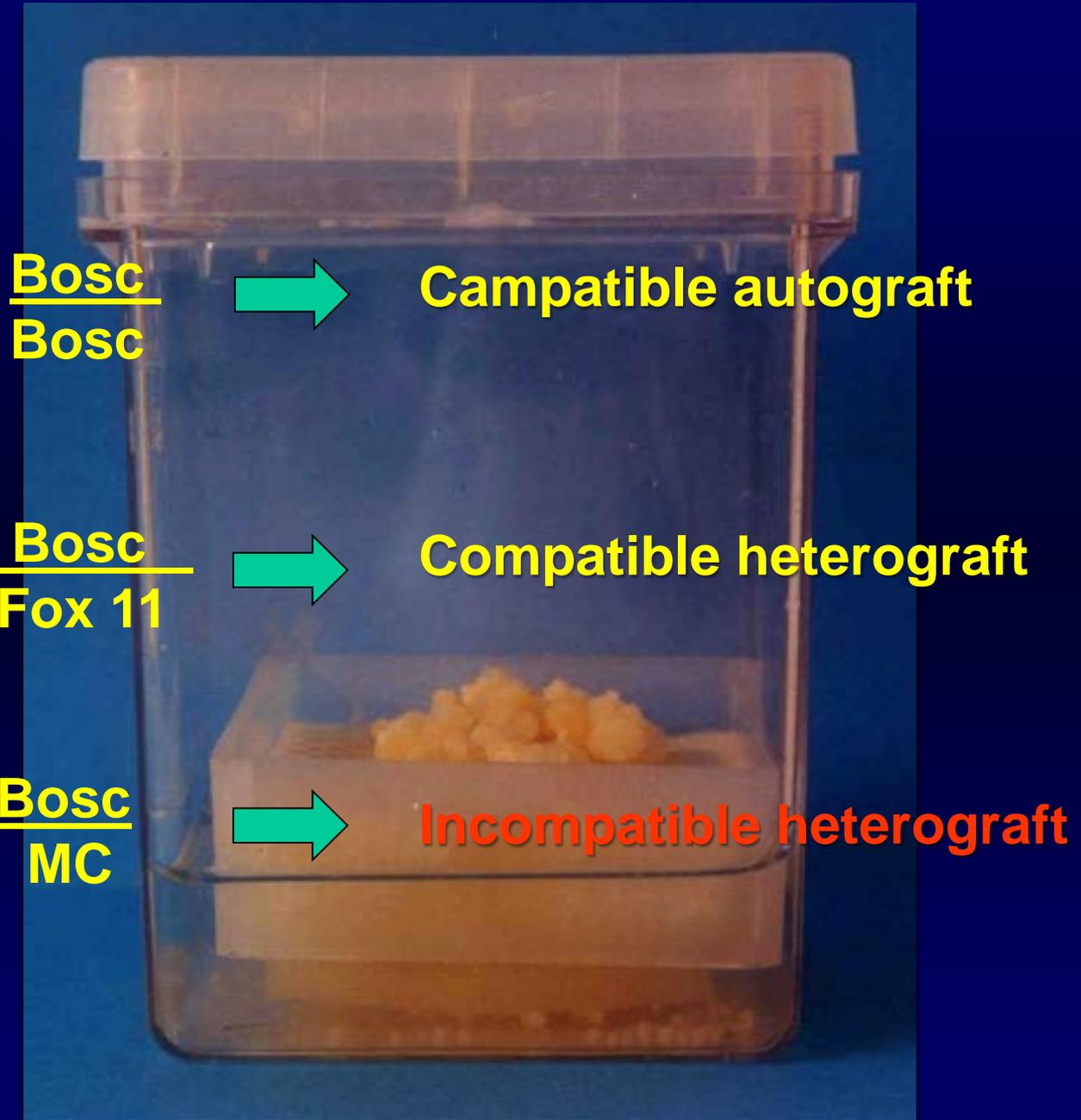


RNA ANALYSES:

mRNA DIFFERENTIAL DISPLAY

allows direct isolation and cloning of cDNA fragments corresponding to mRNAs differently expressed in the compatible and incompatible combinations *(Pirovano et al., 2002)*

- RNA extraction
- Reverse transcription with 3'-anchored primer (T₁₂AC)
- PCR amplification with 3'-anchored primer and random decamer
- Separation on acrylamide gels and identification of DD-fragments
- DNA extraction from excised bands of interest
- Re-amplification by PCR with same conditions
- Separation on agarose gels
- Excision of amplified products and cloning and sequencing



Bosc
Bosc



Compatible autograft

Bosc
Fox 11



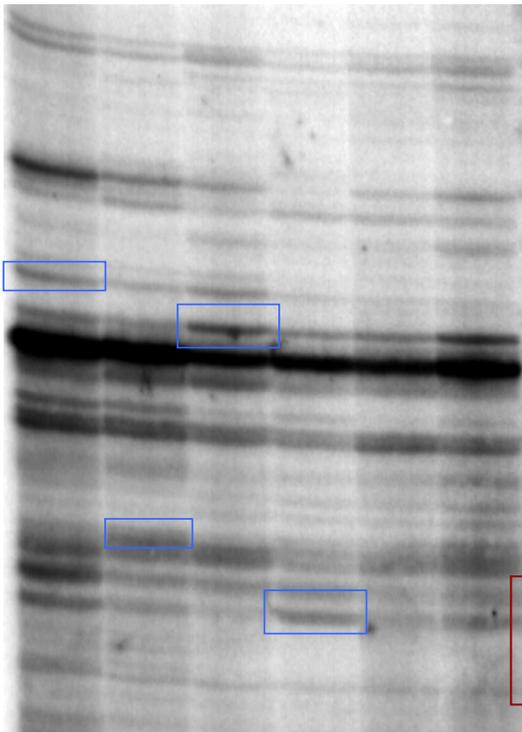
Compatible heterograft

Bosc
MC



Incompatible heterograft

mRNA Differential Display



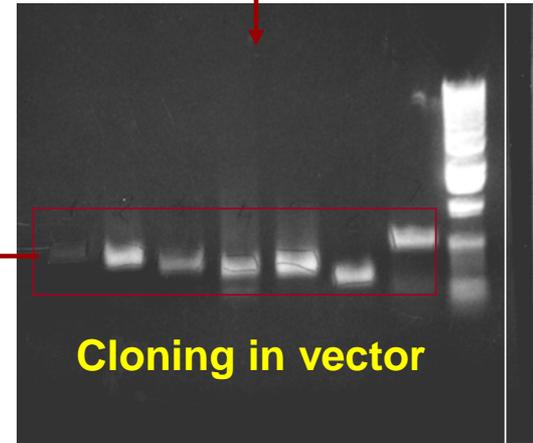
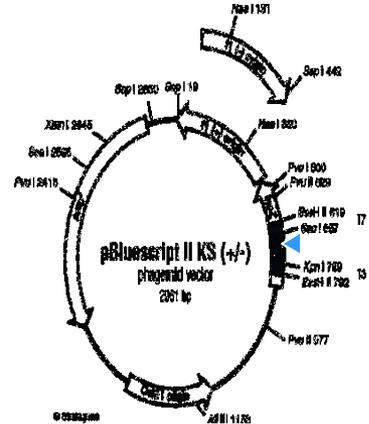
B/B B/E B/F B/B B/E B/F

Primers:

T₁₂AC; T₁₂CA; T₁₂AA; T₁₂GA; T₁₂AG; T₁₂CG

GTGGCCGATG
AGCAGCGAGG
CCTGGGTCAG
GTCGGTTGTC

Reamplification of cDNA fragments



Reverse Northern Dot-Blot
To screen cDNA fragments
really differentially expressed

Northern blot
to study gene expression

Sequencing cDNAs

**Homology research in
GenBank
and EMBL databases**

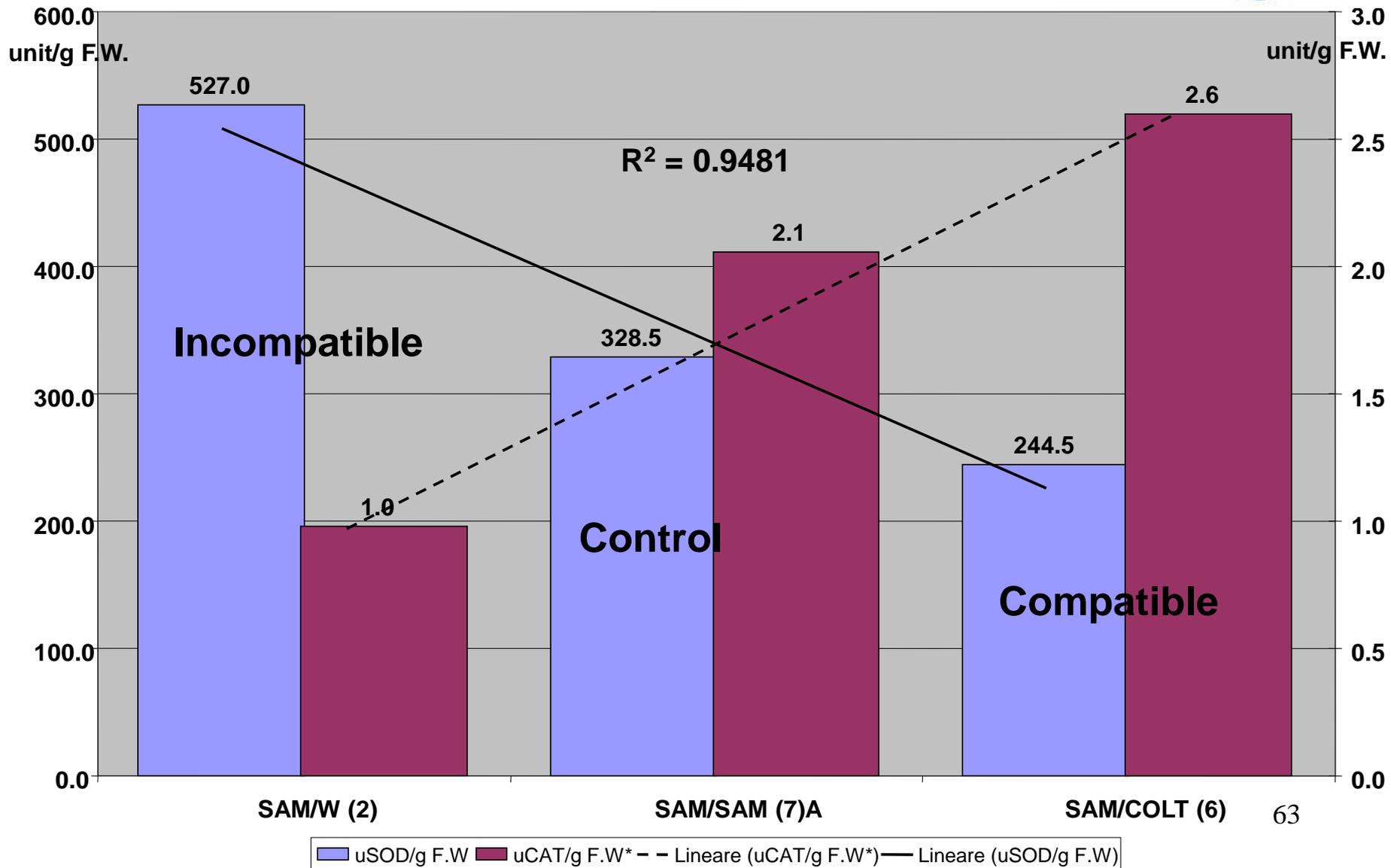
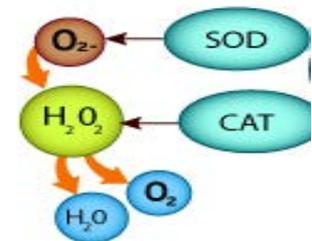
Clone	Tag size (pb)	Putative identity	Associated sequence	% identity, expect value	Northern analysis
T12CA-7	292	NT3	<i>Nicotiana tabacum</i>	39% 5e-10	
T12CA-10	138	Pathogenesis related protein I (SaPRI)	<i>Santalum album</i>	100% 4e-53	
T12AG-14	261	Unknown protein	<i>Arabidopsis thaliana</i>	60% 9e-15	
T12AG-7	215	Import intermediate associated protein (chloroplast)	<i>Pisum sativum</i>	86% 3e-24	
T12AG-10	195	Extracellular dermal glycoprotein (EDGP)	<i>Daucus carota</i>	89% 1e-13	
T12AG-14	383	Putative protein translation factor SUI1 homolog (eIF-2A)	<i>Arabidopsis thaliana</i>	92% 8e-15	
T12CG-7	302	NADH dehydrogenase (ubiquinone) subunit 1	<i>A. Thaliana</i>	94% 2e-20	
T12CG-11	192	Platelet/endothelial cell adhesion molecule	<i>Homo sapiens</i>	61% 5e-3	
T12AA-14	113	No significant similarity			

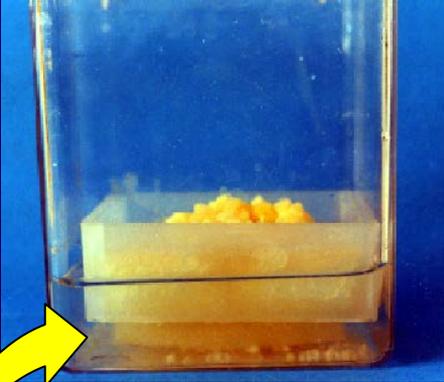
$\frac{B}{B}$ $\frac{B}{MC}$ $\frac{B}{F}$

PROTEINS ANALYSES

- **Enzyme activity (SOD and CAT)**
- **SDS-PAGE** (sodium dodecyl sulfate polyacrylamide gel electrophoresis)
- **2D-PAGE** (Two-dimensional gel electrophoresis)

SOD and CAT activities in cherry co-culture





**Proteins
extracted
from the
liquid
medium**

1 - MC flask

2- Bosc/MC

3 - BH/MC

kDa

200

116

97

66

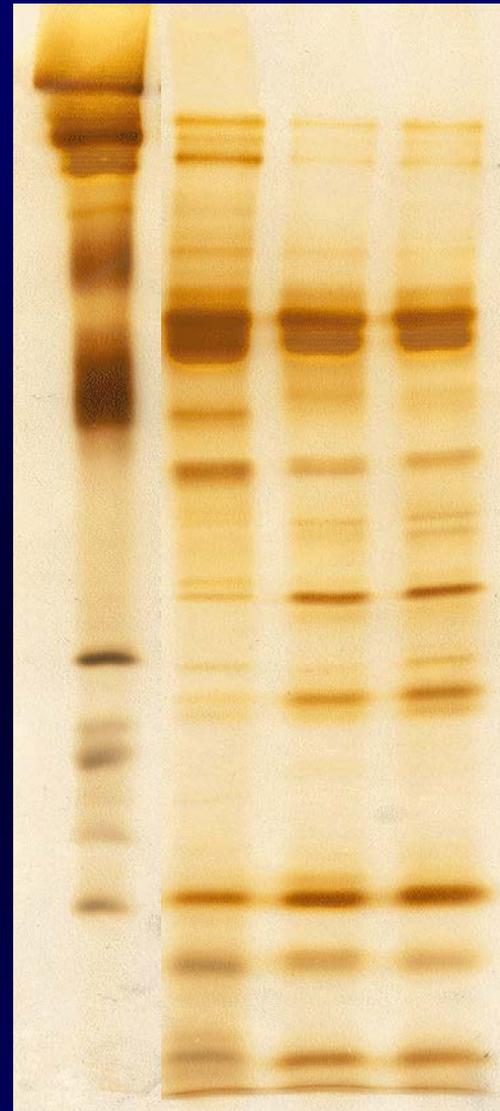
45

31

21

14

6



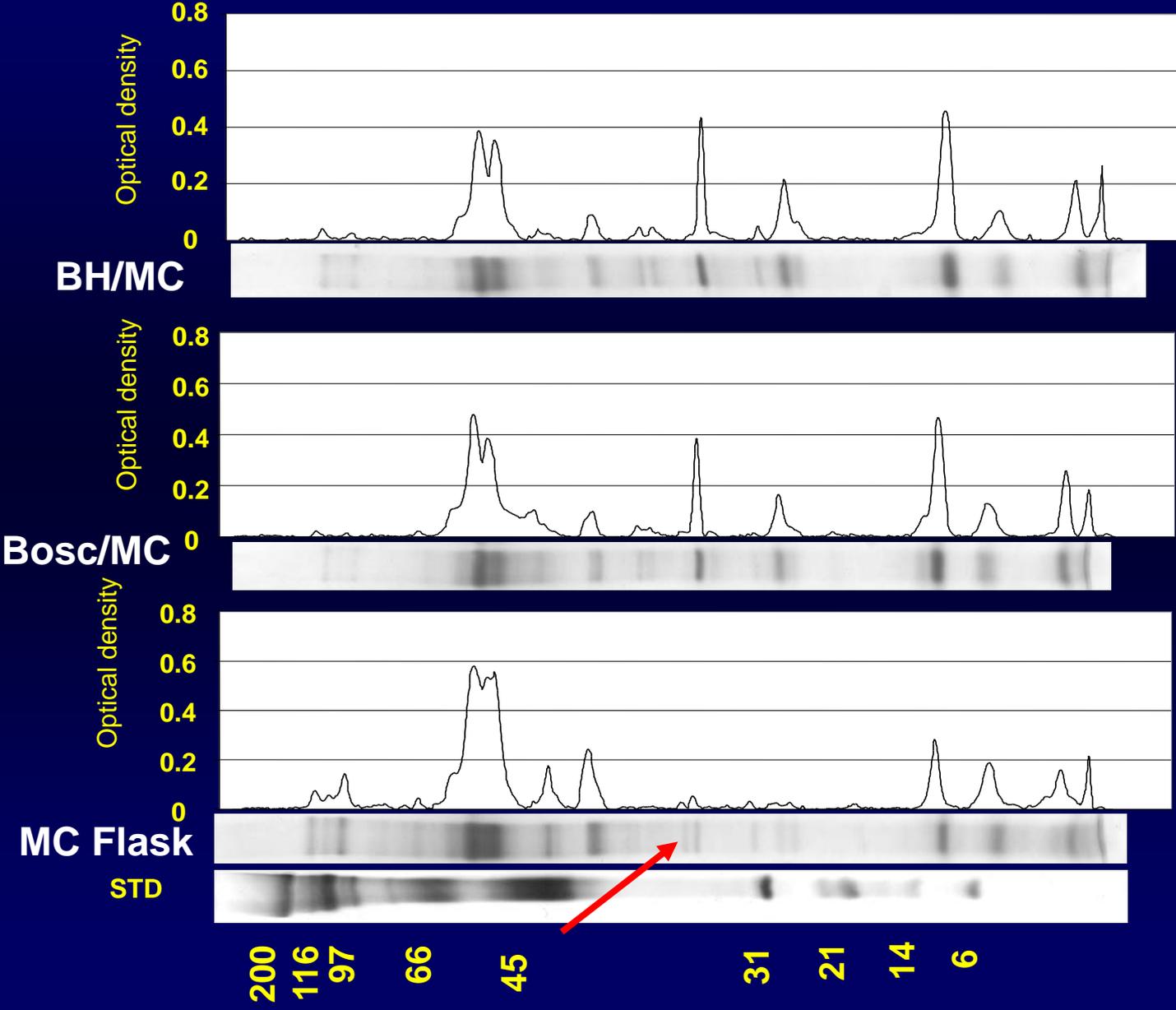
STD

MC flask

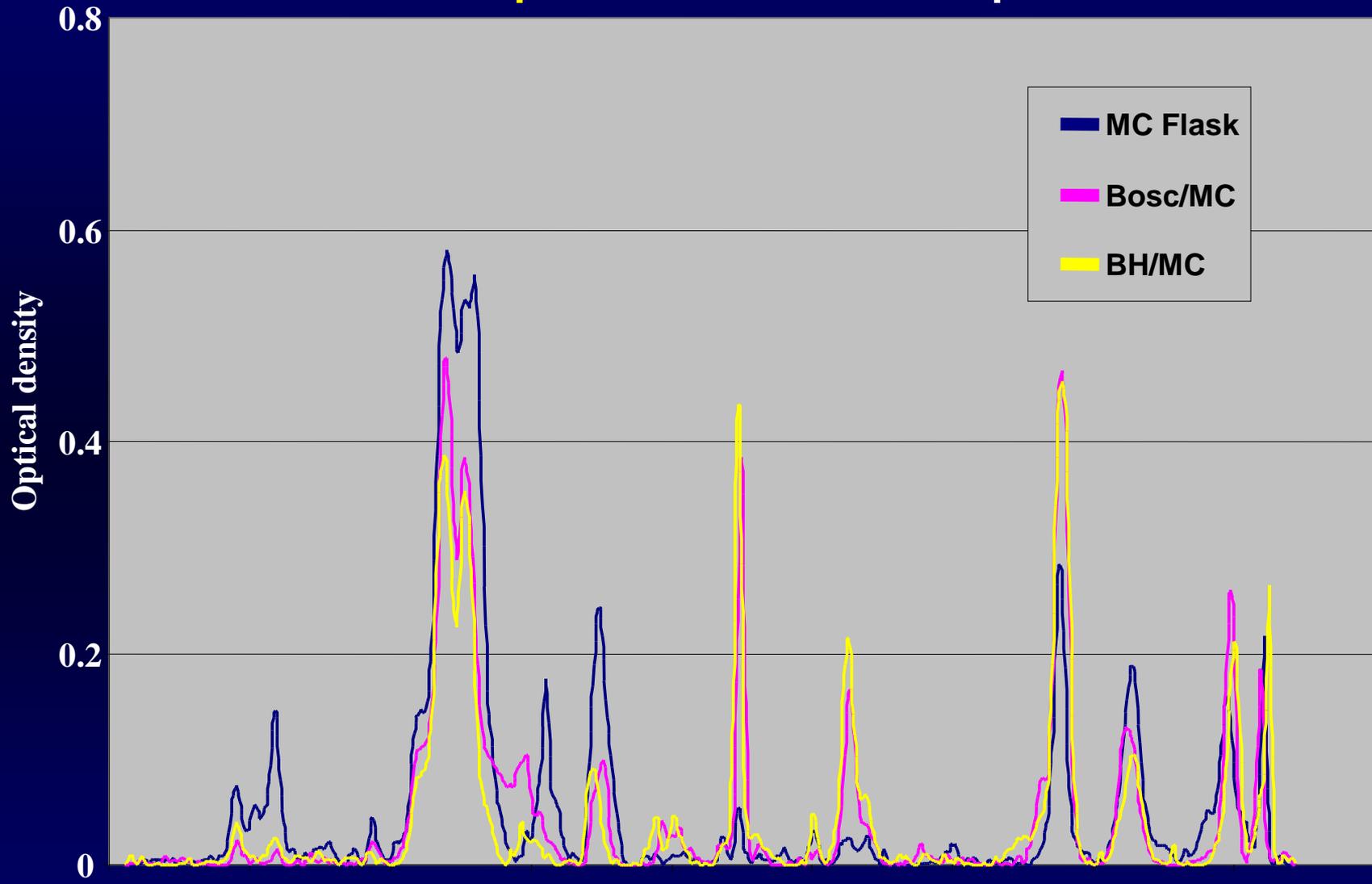
BOSC/MC

BH/MC

Quince MC: densitometric analysis of proteins extracted from the liquid medium of cell suspension alone and in co-culture



Quince MC: densitometric analysis of proteins extracted from the liquid medium of cell suspension



STD



kDa

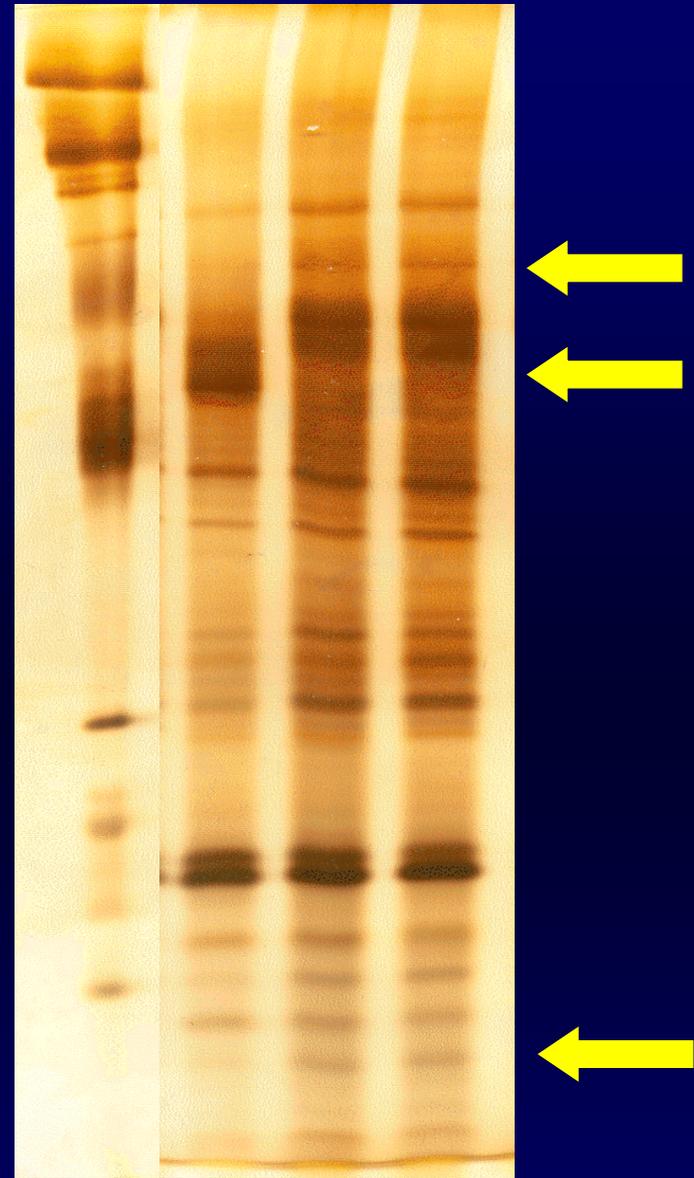
200 116 97 66 45 31 21 14 6



Proteins
extracted from
callus above
the porous
membrane

1. **Bosc/Bosc**
2. **Bosc/Fox 11**
3. **Bosc/MC**

kDa
200
116
97
66
45
31
21
14
6



STD

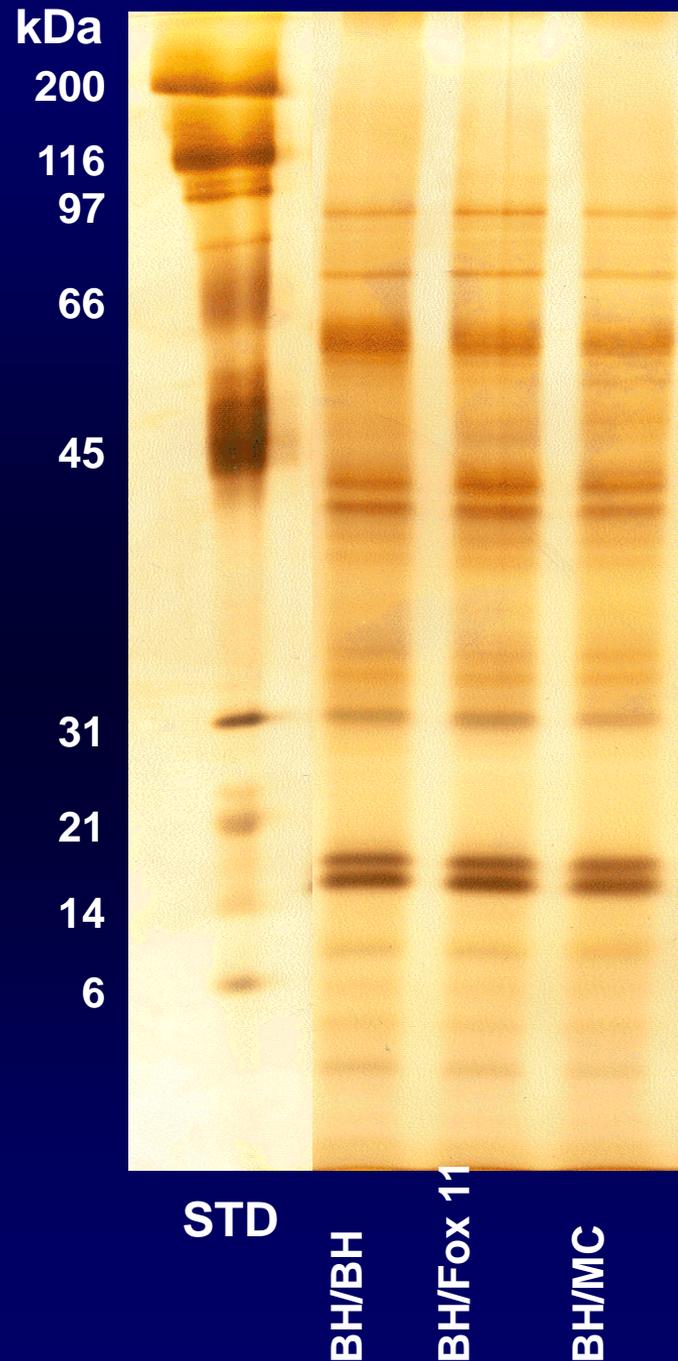
Bosc/Bosc

Bosc/Fox 11

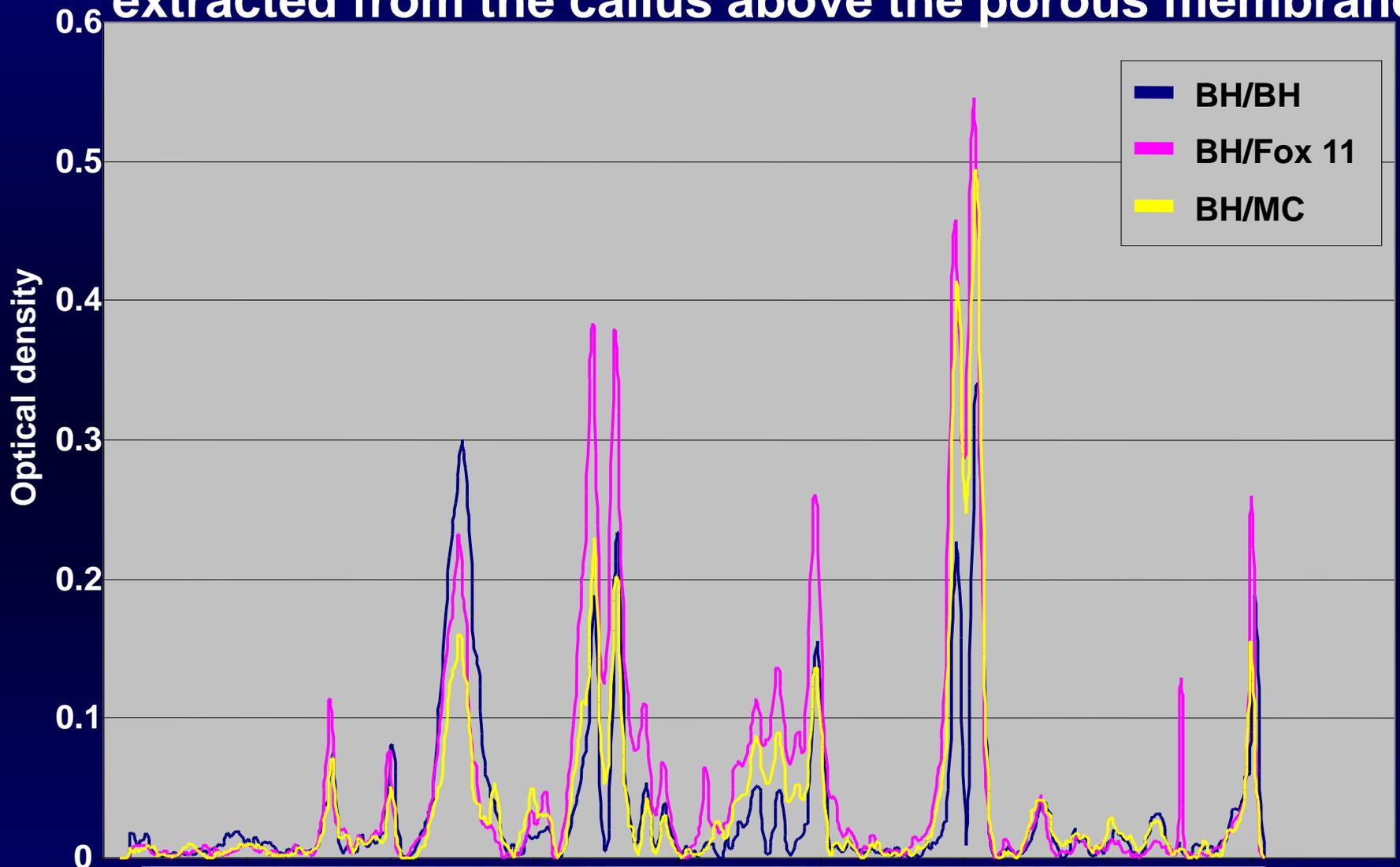
Bosc/MC

Proteins
extracted
from **callus**
above the
porous
membrane

1. **BH/BH**
2. **BH/Fox 11**
3. **BH/MC**



Cv. Beurré Hardy: densitometric analysis of proteins extracted from the callus above the porous membrane



STD

200

116

97

66

45

31

21

14

6

2D-PAGE



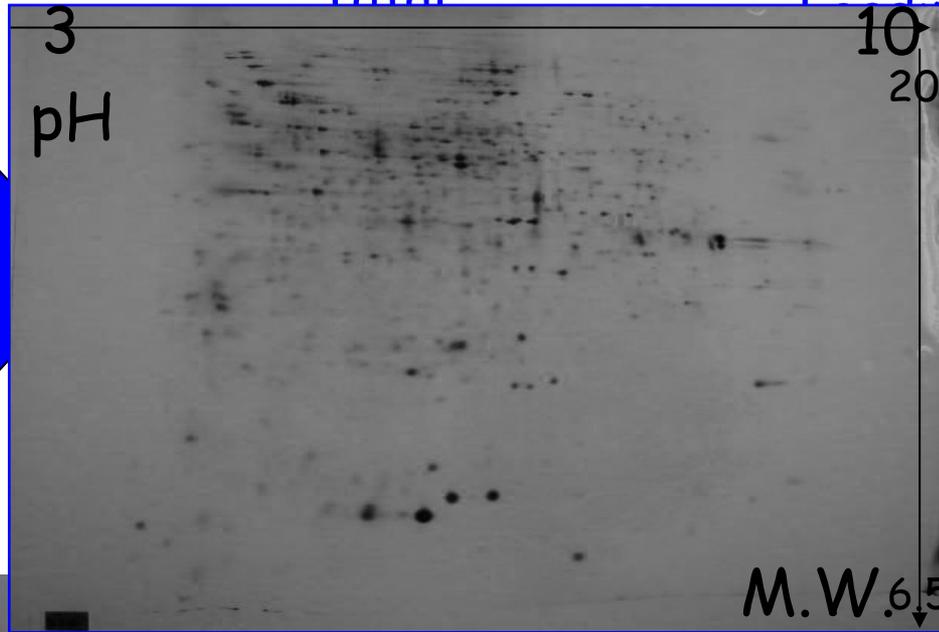
Starting material



Total



Loading of first dimension



First dimension

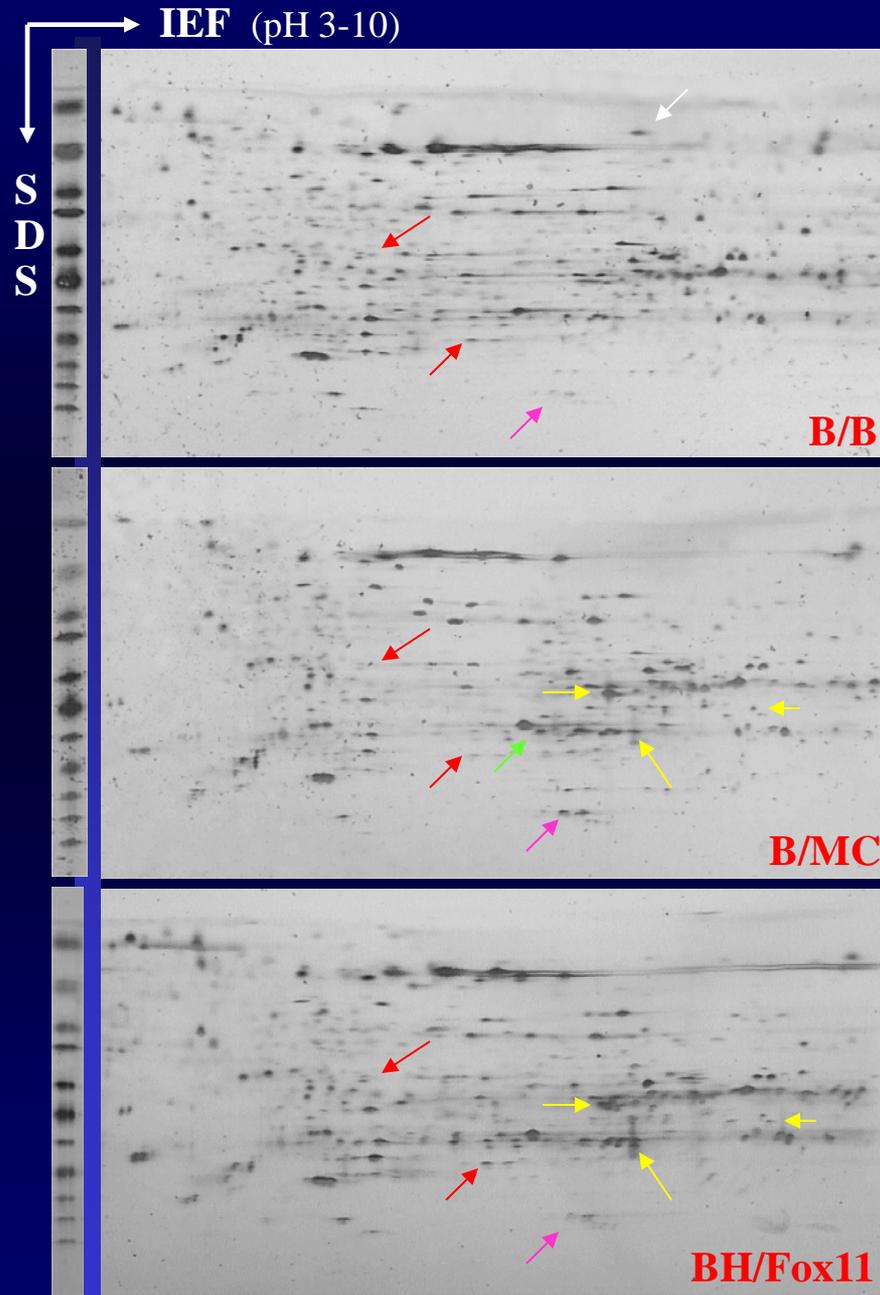


Staining



Second dimension

Protein expression by 2D electrophoresis



only in B/B

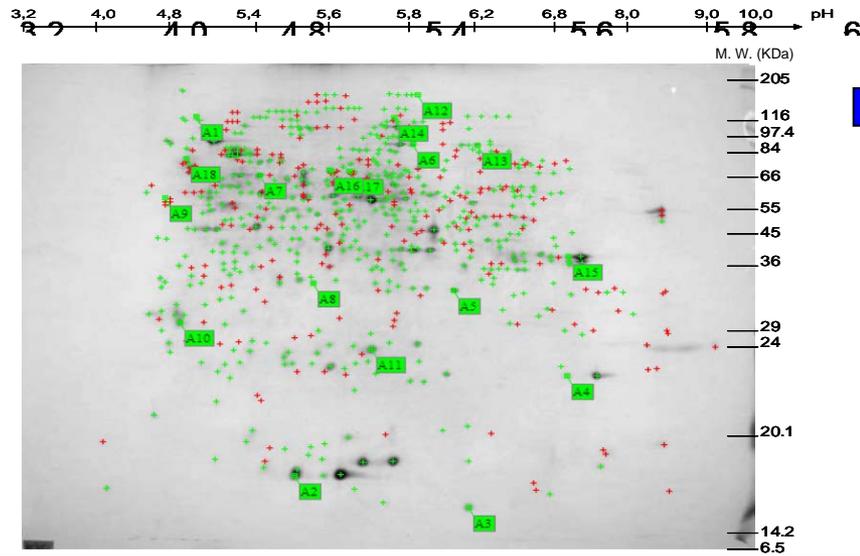
only in B/MC

decrease in B/MC

increase in BMC

only in B/MC and BH/Fox11

Gel comparison between pear combinations

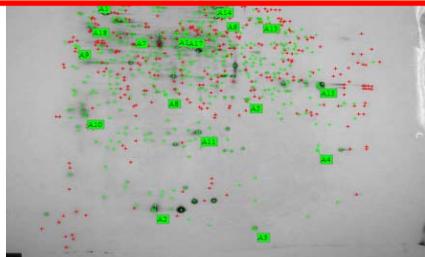


n° spots: 706

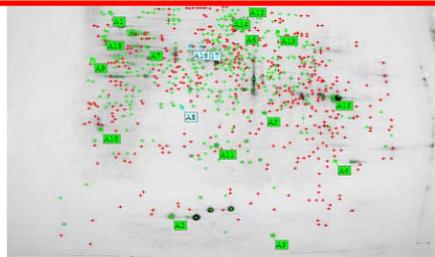
n° spots: 802
 matches: 476
 % matches: 63%

The incompatible combination exhibits the gel with the higher spots number

n° spots: 857
 matches: 506
 % matches: 65%



K/K

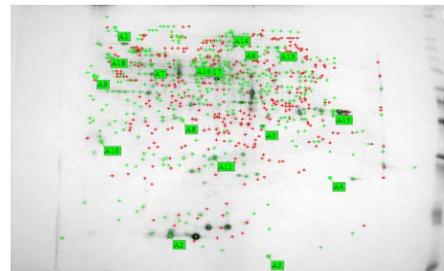


K/MC

n° spots: 774

matches: 425

% matches: 57%



K/A28



Starting from Pirovano *et al.* (2002) we tried to identify the following proteins in a databank.

Protein	Species	Homology level	Expression level
NT3	<i>Nicotiana tabacum</i>	39%	
Patogenesis Related Protein I (SaPRI)	<i>Santalum album</i>	100%	
Import intermediate associated protein	<i>Pisum sativum</i>	86%	
Extracellular dermal glycoprotein (EDGP)	<i>Daucus carota</i>	89%	
Protein translation factor SUI1 homolog (eIF-2A)	<i>Arabidopsis thaliana</i>	92%	
NADH dehydrogenase Subunit I	<i>Arabidopsis thaliana</i>	94%	
Platelet endothelial cell adhesion molecule	<i>Homo sapiens</i>	61%	

K K₇₄ K
K MC F

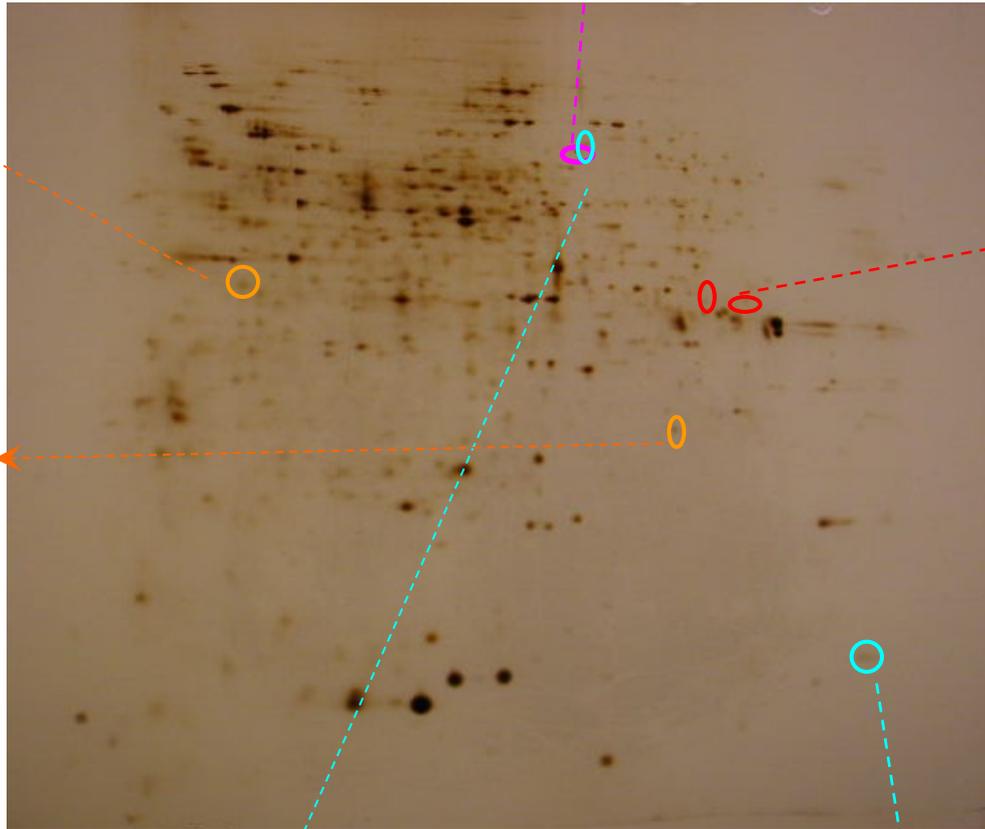
Protein name	Accession number	Entry name	Molecular weight (KDa)	Theoretical pI
NT3	Q9XEY9	Q9XEY9_TOBAC	70.78	-
hrgpNT3	P13983	EXTN_TOBAC	65.41	10.00
Pathogenesis Related Protein I (SaPRI)	O22479	O22479_9MAGN	15.31	-
Import intermediate associated protein	Q43715	TOC75_PEA	88.27	7.00
Extracellular dermal glycoprotein (EDGP)	Q39688	EPIG_DAUCA	43.55	7.97
Protein translation factor SUII homolog (eIF-2A)	Q9FE78	Q9FE78_ARATH	38.80	5.00
Protein translation factor SUII homolog (eIF-2A) (subunit β)	Q41969	IF2B_ARATH	30.07	6.79
Putative NADH dehydrogenase Subunit I	Q9M9M9	N7BM_ARATH	18.31	9.26
NADH dehydrogenase Subunit I (75 Kda subunit)	Q9FG16	NUAM_ARATH	81.50	6.24
NADH dehydrogenase Subunit I (18 Kda subunit)	Q9FLX7	NUFM_ARATH	19.18	4.73
Platelet endothelial cell adhesion molecule	P16284	PECA1_HUMAN	82.53	6.55

Grid references

Import intermediate associated protein

eIF-2A

Extracellular dermal glycoprotein (EDGP)



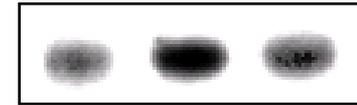
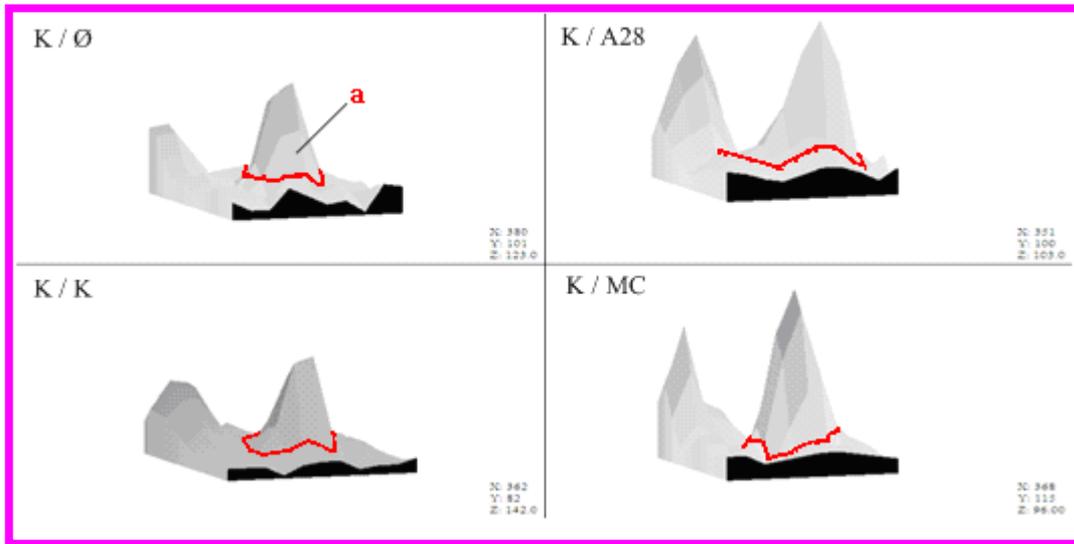
Subunità β dell'eIF-2A

Putative NADH dehydrogenase

Subunità I della NADH dehydrogenase

For these proteins, a correlation between transcripts and spot intensities have been found.

Import intermediate associated protein

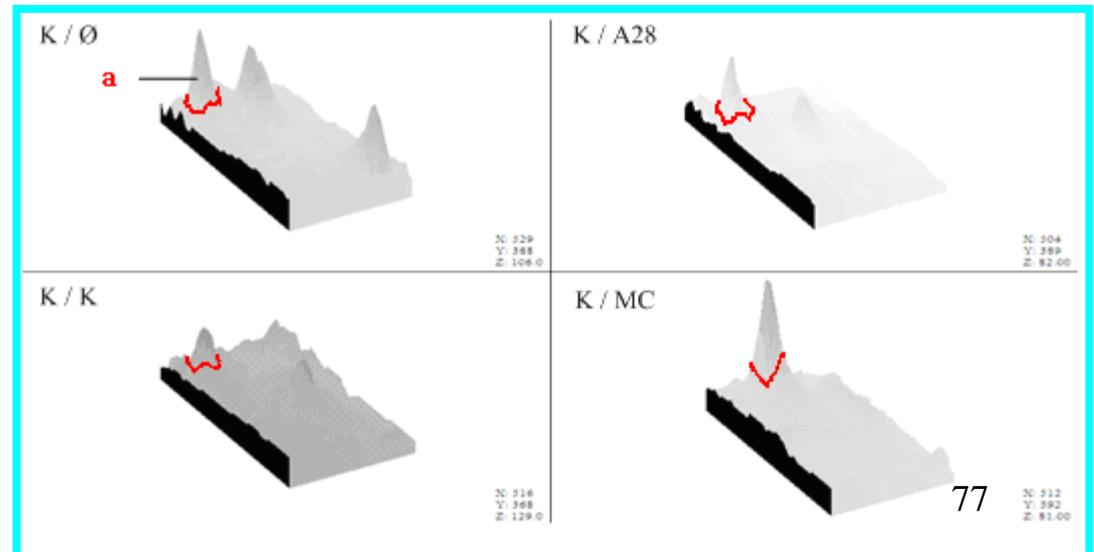


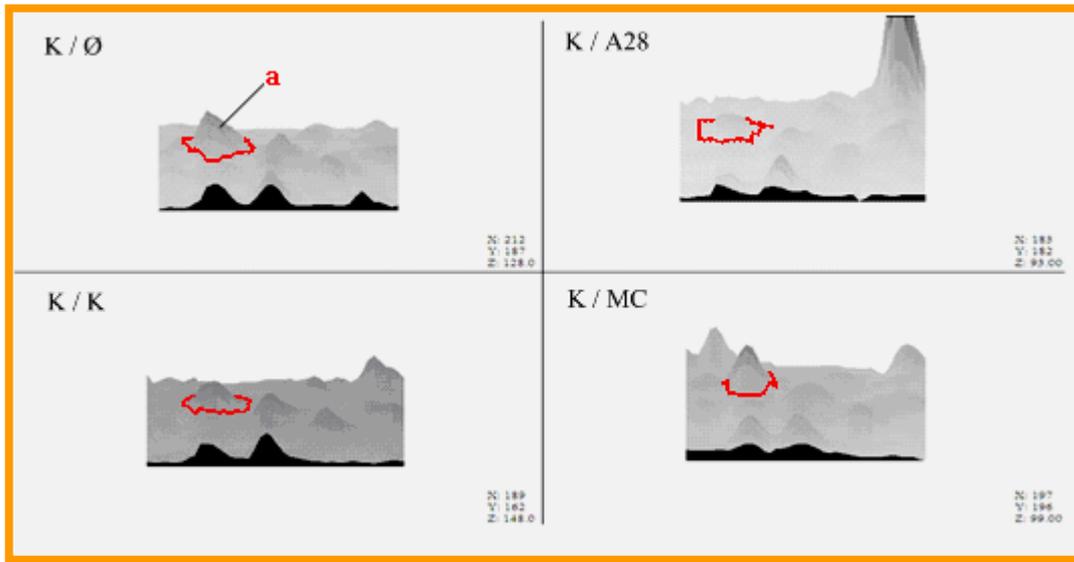
K K K
K MC A28

Putative NADH dehydrogenase

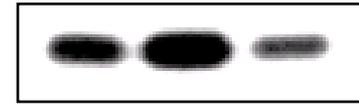


K K K
K MC A28





eIF-2A

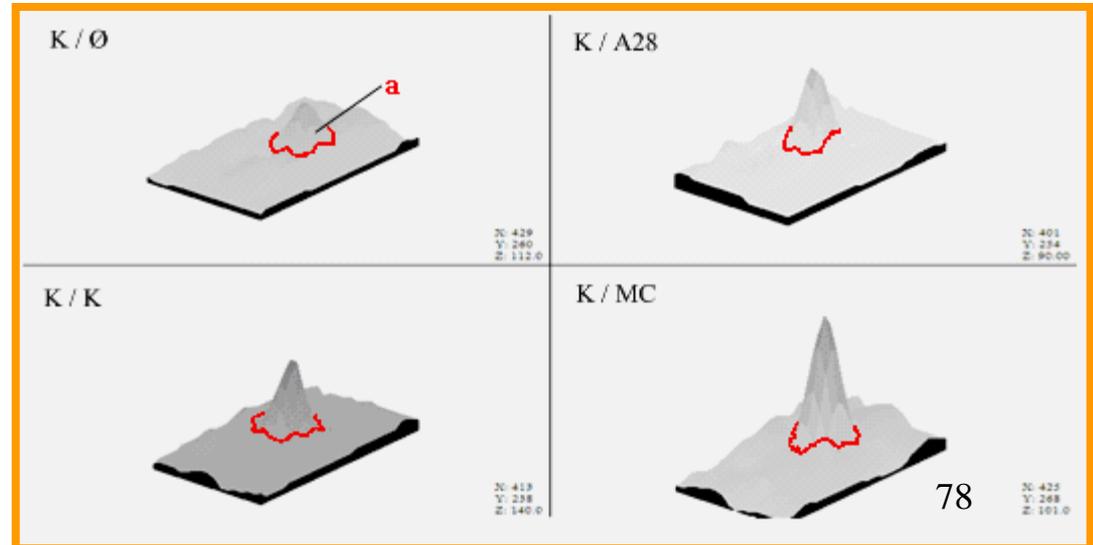


 K K K
 ———
 K MC A28

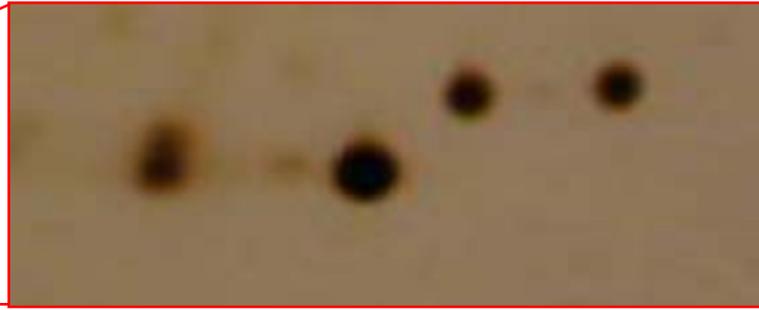
Subunit β of IF-2A



 K K K
 ———
 K MC A28

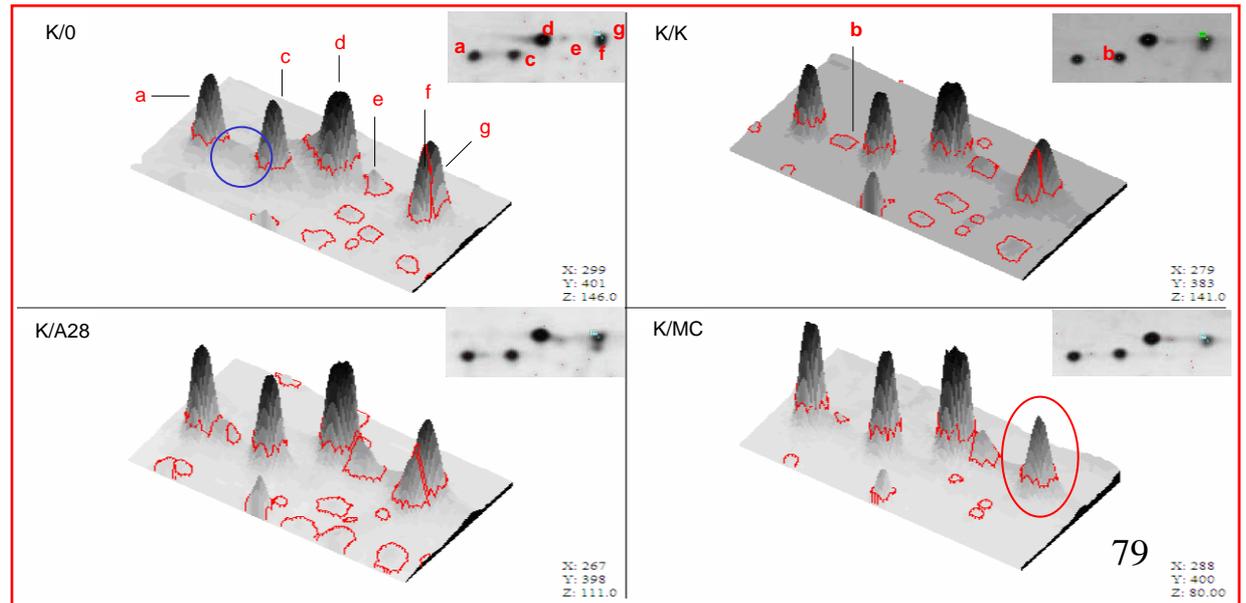
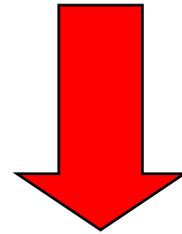


Gel analysis

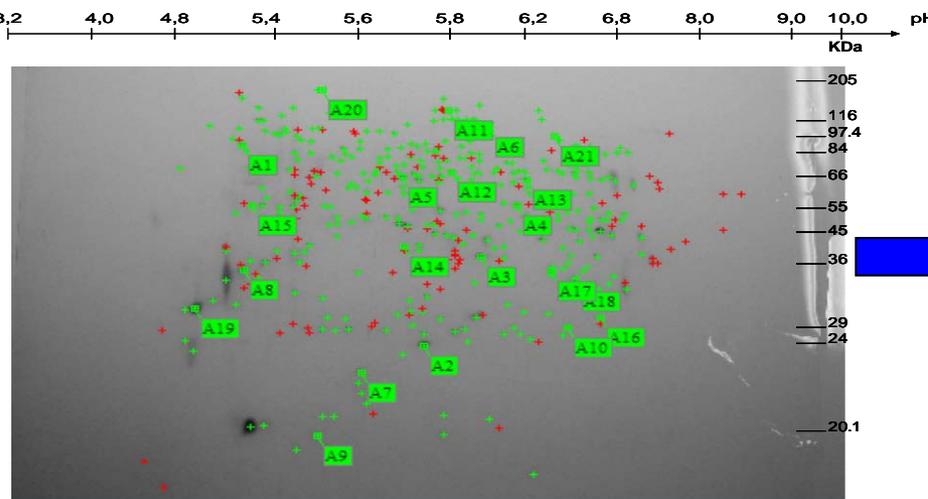


KAISER/KAISER

Details



Gel comparison between cherry combinations

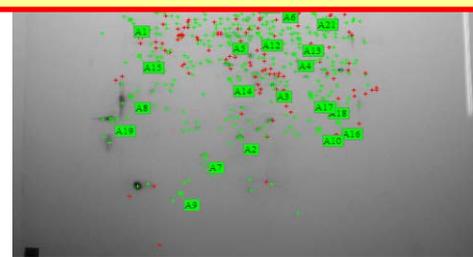


n° spots: 435

- The compatible combination exhibits the gel with the highest number of spots.
- The combination with the highest level of homology is the self-graft.
- The values of percent matches decrease with the increase of the graft-incompatibility level.

n° spots
match
% ma

: 451
: 312
: 70%



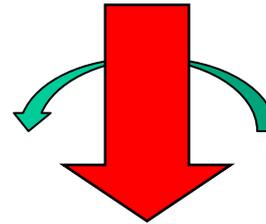
n° spots: 471
matches: 326
% matches: 72%⁸⁰

S/C

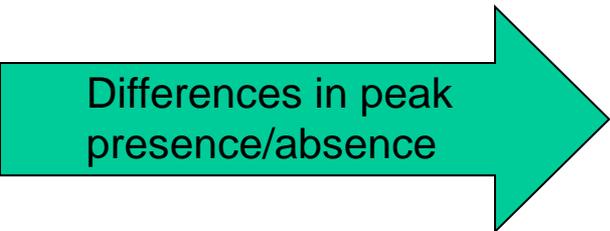
Gel analysis



SAM/SAM

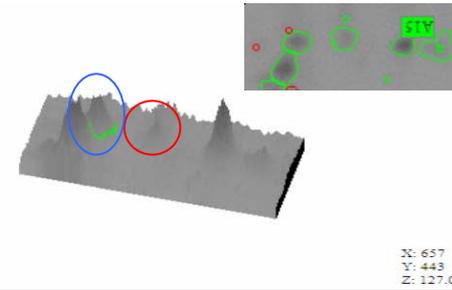


Spot area



S / Ø

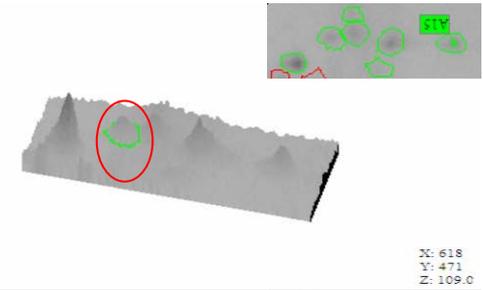
A



X: 657
Y: 443
Z: 127.0

S / C

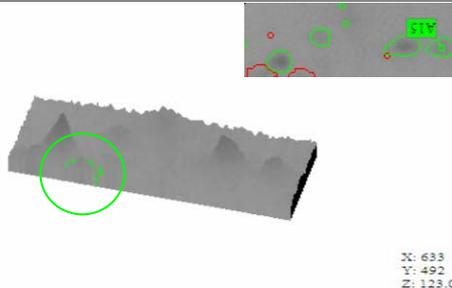
B



X: 618
Y: 471
Z: 109.0

S / W

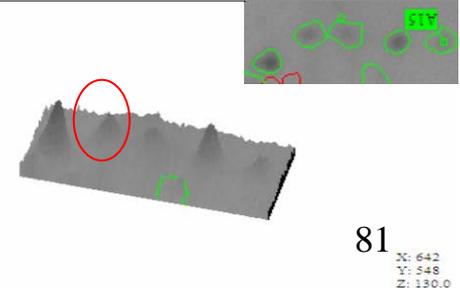
C



X: 633
Y: 492
Z: 123.0

S / S

D



81

X: 642
Y: 548
Z: 130.0

CONCLUSIONS

- The “*in vitro*” model provides a lot of information about relationship between genotypes and cell-cell recognition
- Beurré Bosc showed a **higher growth** compared to Beurré Hardy
- **Callus respiration** of Bosc seems to be higher in presence of incompatible rootstock quince C.
- **Proteins changes** can be associated with cell-cell recognition mechanism.
- Identification of proteins as ‘messengers’ of biological cell-cell recognition between the two genotypes in incompatible graft combination would provide an **early screening “marker”**.

Conclusions

- Gene expression is affected by callus co-culture combination
- The effect on gene expression does not require direct tissue contact: *is a mobile solute involved?*
- Some differentially expressed cDNAs match with nucleotide sequences coding for proteins involved in: **cell adhesion** (T₁₂CG11), **senescence** and/or **programmed cell death** (T₁₂CA-10; T₁₂AG-14), **wounding response** (T₁₂AG-10) and **tracheary element differentiation** (T₁₂CG-7).

Acknowledgements

Sara Serra – Washington State University

Vincenzo Ancarani - University of Bologna

Andrea Masia - University of Bologna

Gian Attilio Sacchi - University of Milan

Luca Espen - University of Milan

Fabio Nocito - University of Milan

Andrea Fabbri - University of Parma

**THANK YOU FOR YOUR
ATTENTION**