

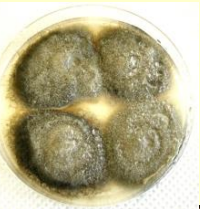


Assimilate partitioning in higher plants and Plant Phenomics

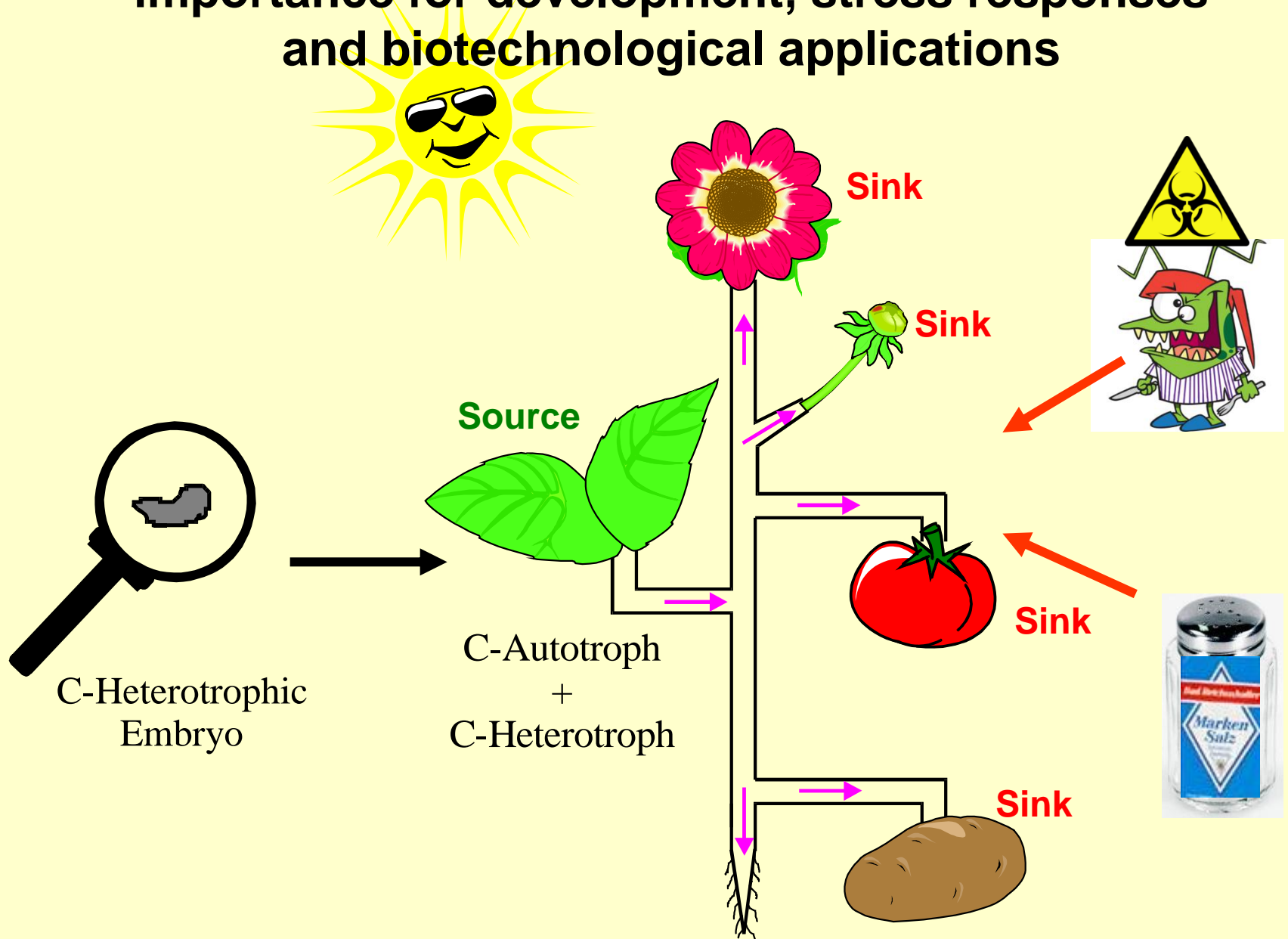
Eric van der Graaff and Thomas Roitsch

University of Copenhagen

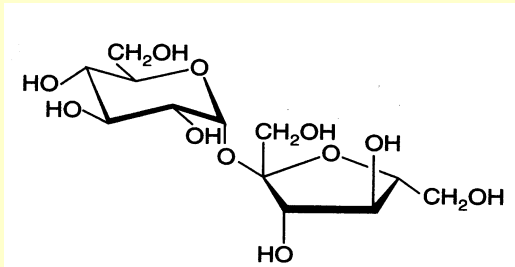
Department of Plant and Environmental Sciences



Assimilate partitioning in higher plants: importance for development, stress responses and biotechnological applications



The Invertase Isoenzymes



Sucrose

α -D-Glucopyranosyl- β -D-fructofuranoside



Glucose + Fructose

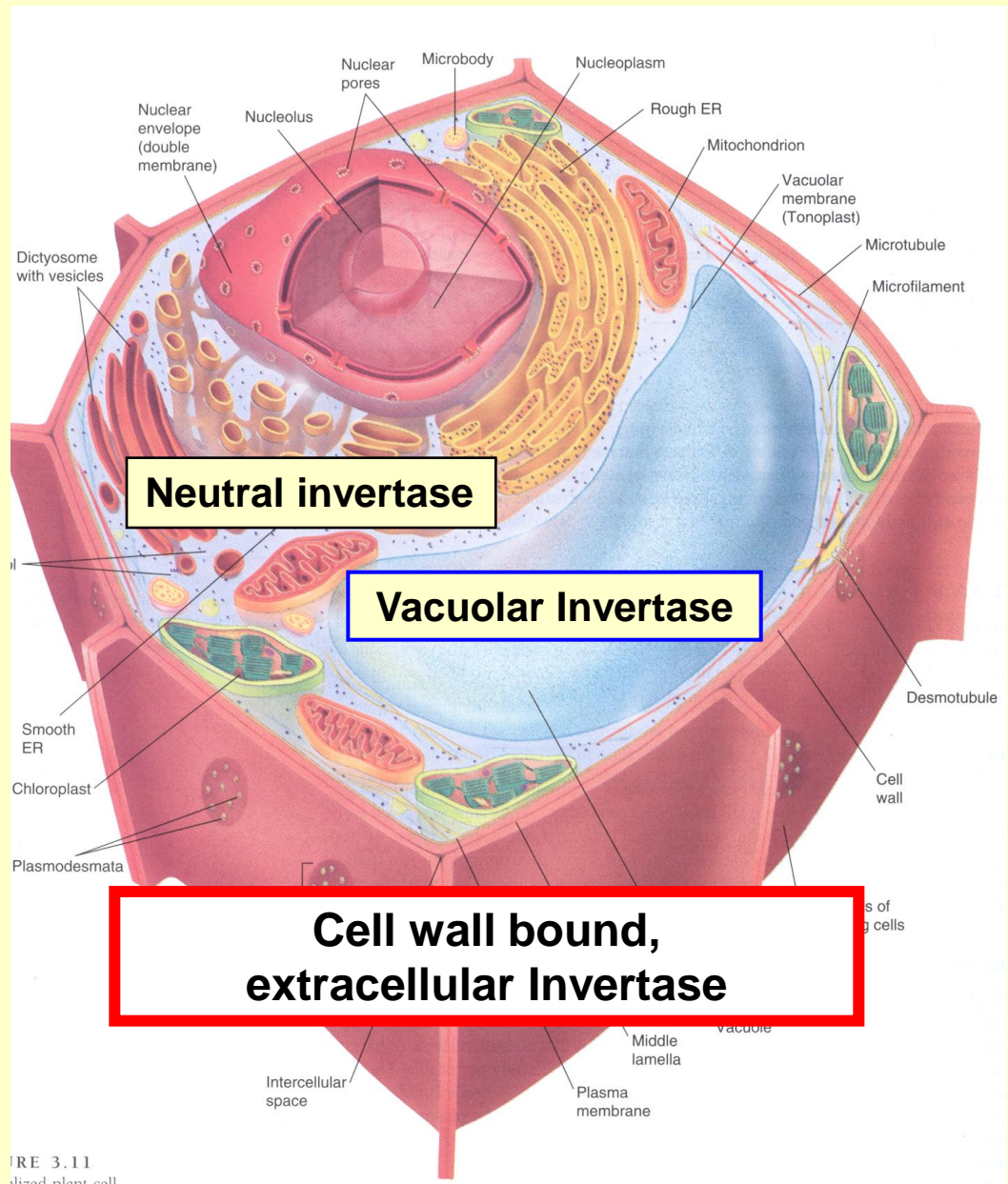
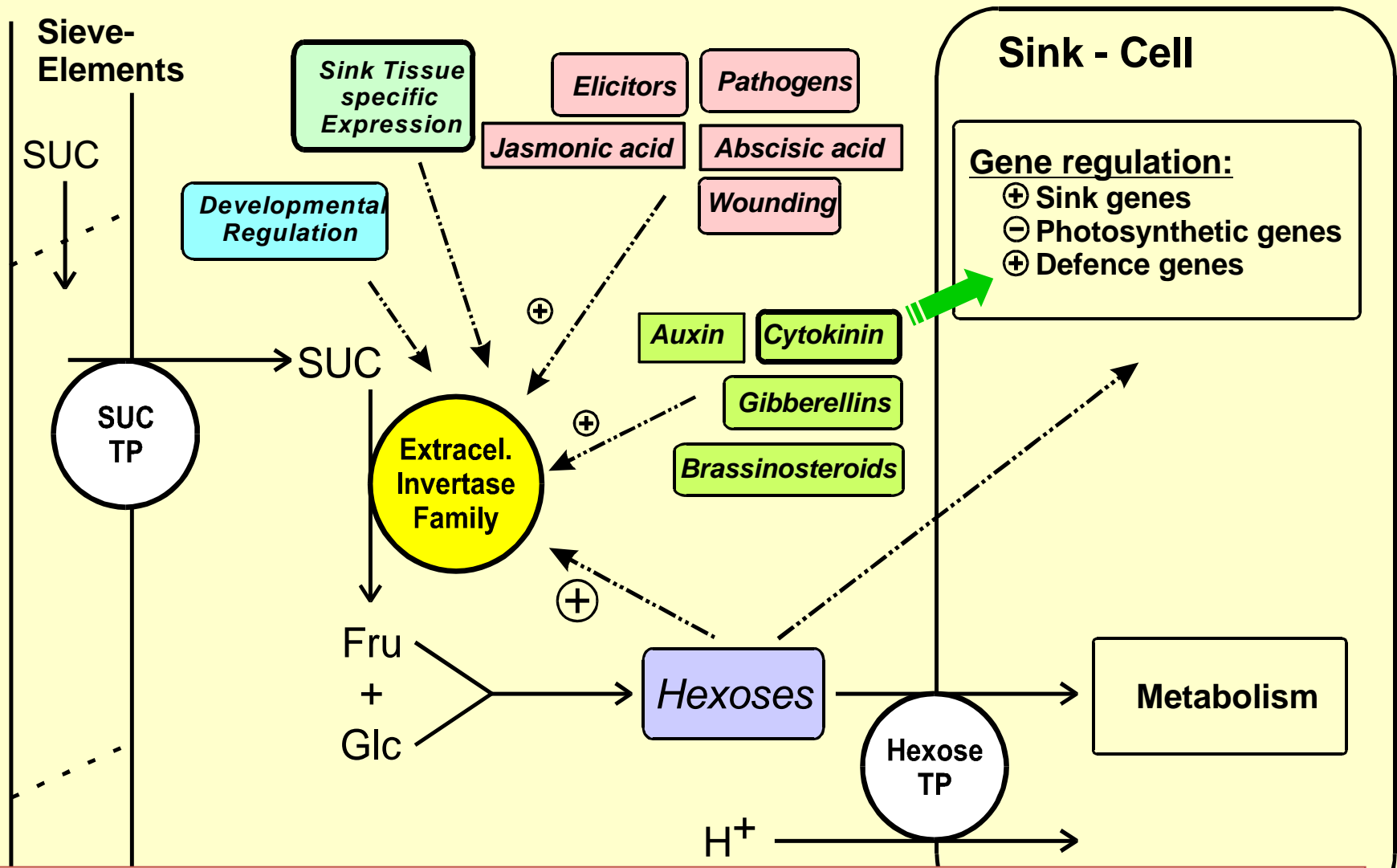


FIGURE 3.11
Invertase in a plant cell.

Extracellular Invertase: Key metabolic Enzyme and PR Protein



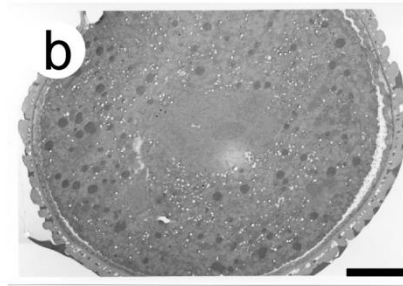
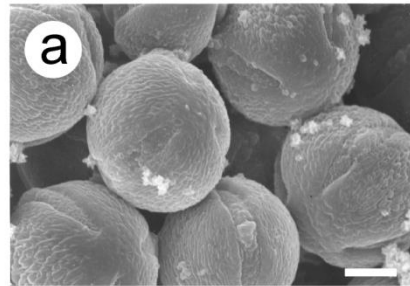
Roitsch (1999) COPS 2, 98f; Roitsch et al. (2004) TIPS 9, 607f; Berger et al. (2007) JXB 58, 4019f
 Albacete et al. (2011) Phyton 50,181; Großkinsky et al. (2012) Plant Sci. 195, 54f

Extracellular invertase is essential for pollen development

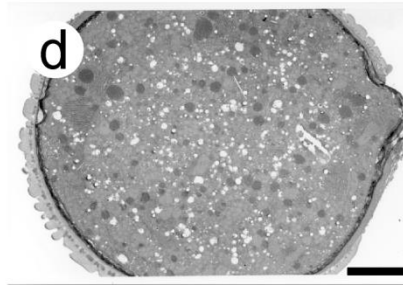
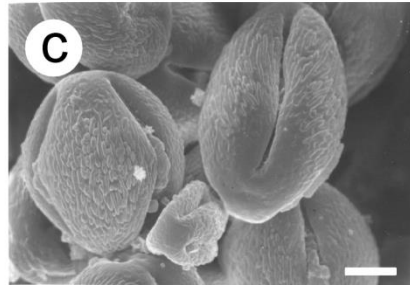
Antisense-Repression of Extracellular Invertase Nin88
results in an Arrest of Pollen Development and male sterile Plants

SEM (2700x)

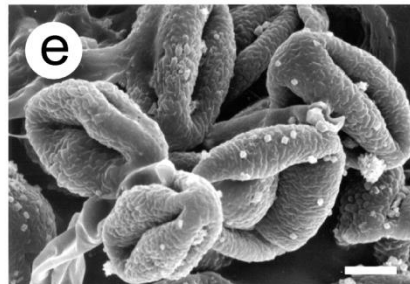
TEM (5600x)



Wildtype

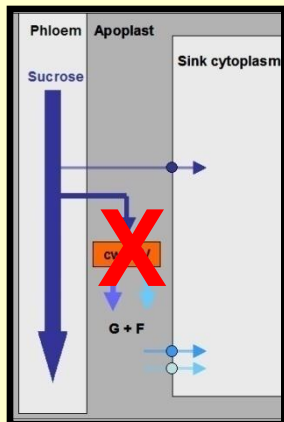


NT23-17



NT23-59

***Nin88::
Nin88-Antisense***



Relation between phytohormone action and carbohydrate metabolism

Extracellular Invertase is essential for the delay of Senescence by cytokinins

Cytokinin → Extracellular Invertase → Delay of Senescence

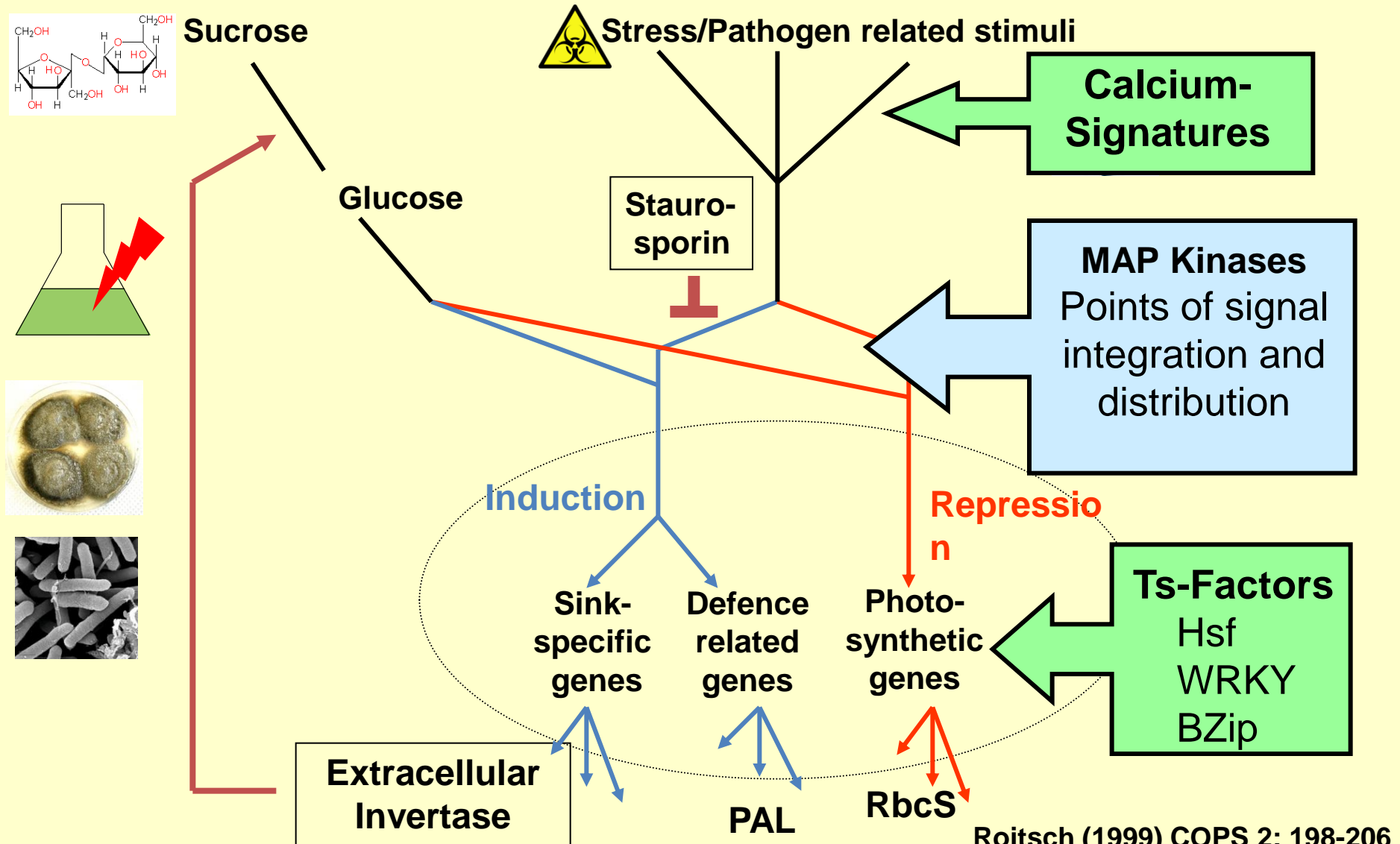
SAG12:
Cin1



W 38

Balibrea et al..
(2004) Plant Cell

Co-ordinated regulation of source/sink relations and defence response by sugars and stress related stimuli



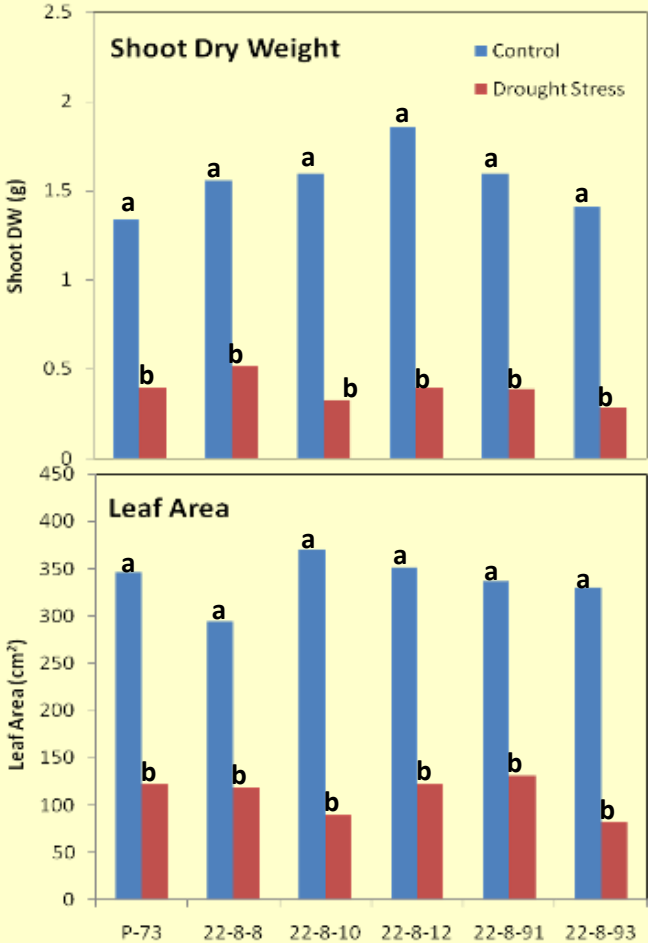
Ectopic expression of Extracellular invertase results in extreme drought stress tolerance

Drought stress tolerance in *CIN1* is not accompanied by a yield penalty under control conditions



WT

CIN1 1-91



Cytokinins stimulate immunity via stimulation of phytoalexin synthesis within the Phytohormones Networks

Phytohormone defence backbone:

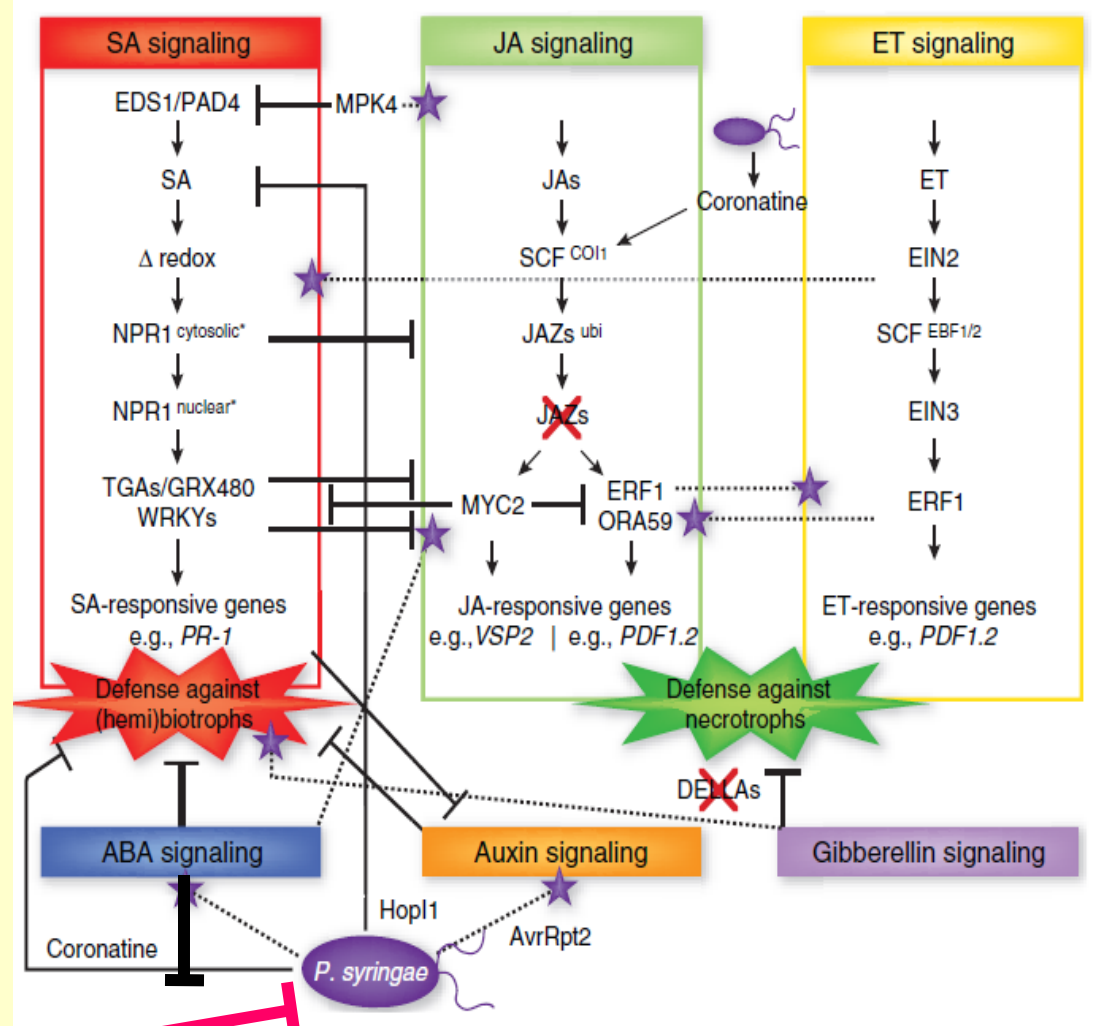
- Salicylic Acid
- Jasmonic Acid
- Ethylene

Additional interactions :

- Abscisic Acid
- Auxin
- Gibberellin

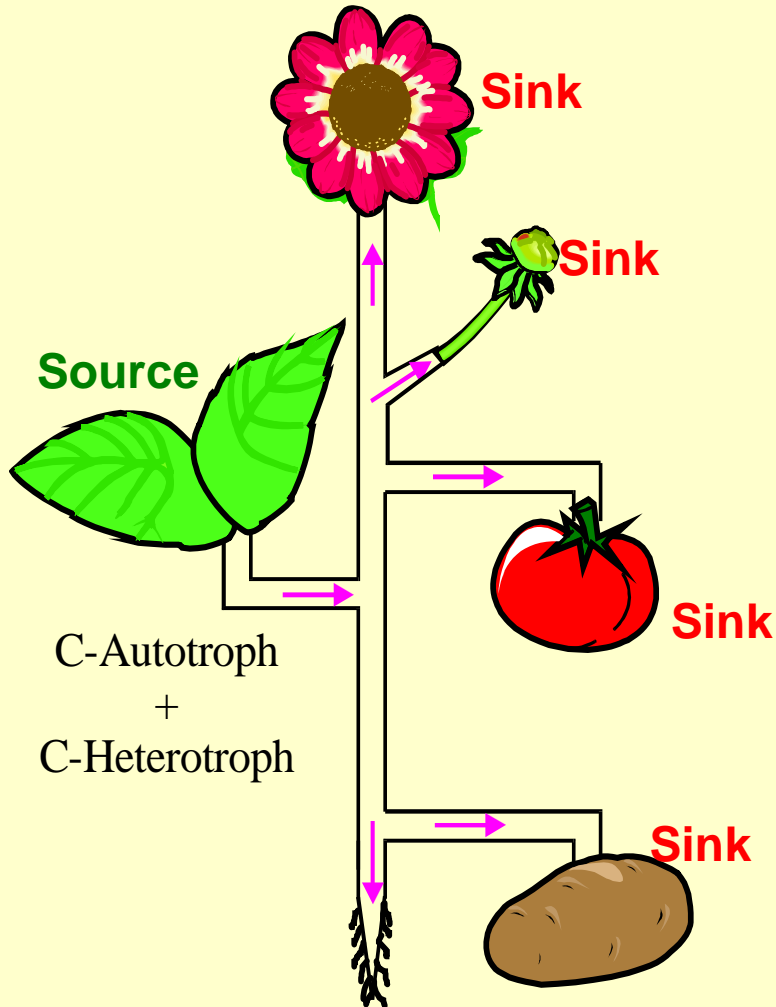


+ Cytokinin signalling
Phytoalexins

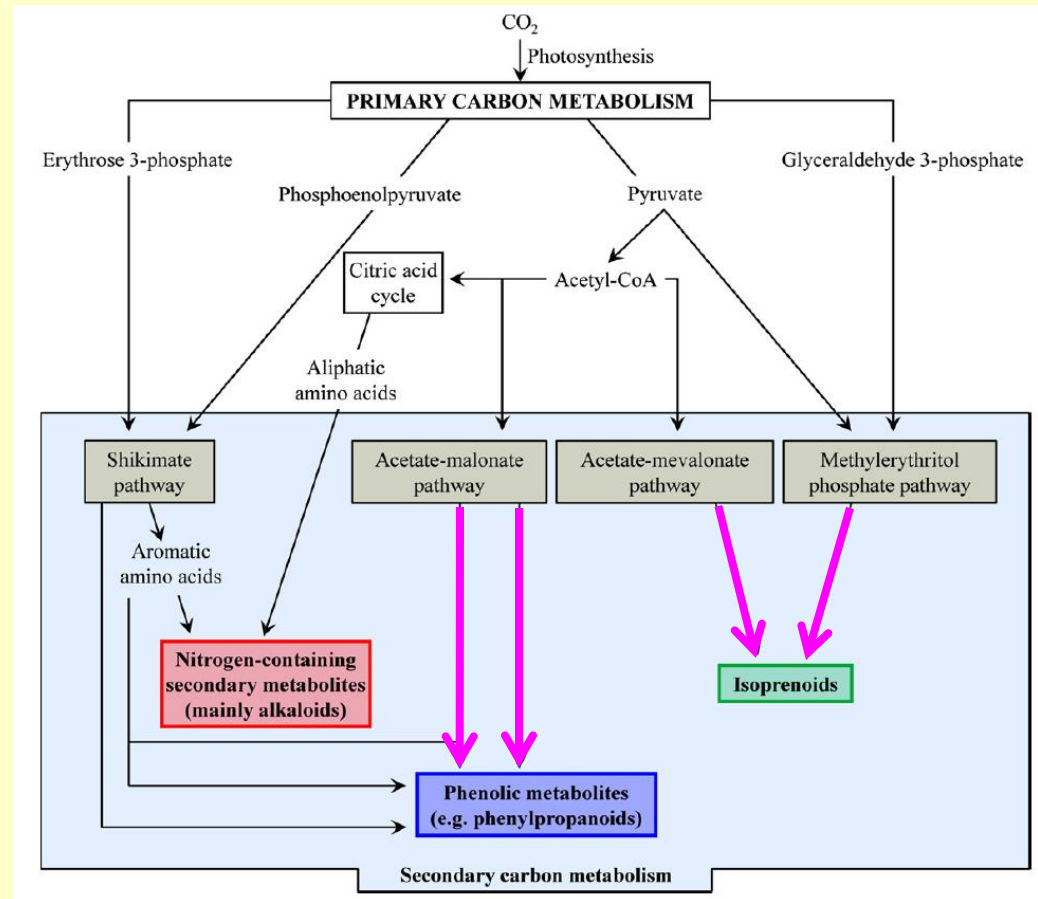


Different scales of carbohydrate partitioning

(A) Long distance transport from source and sink tissues



(B) Metabolic channeling from primary to secondary metabolism

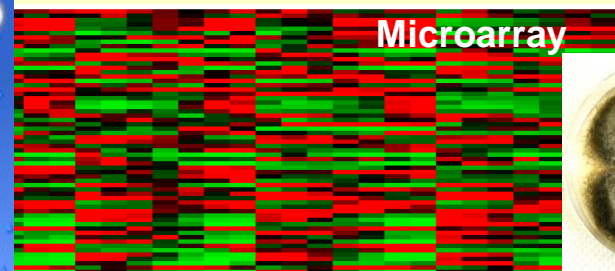
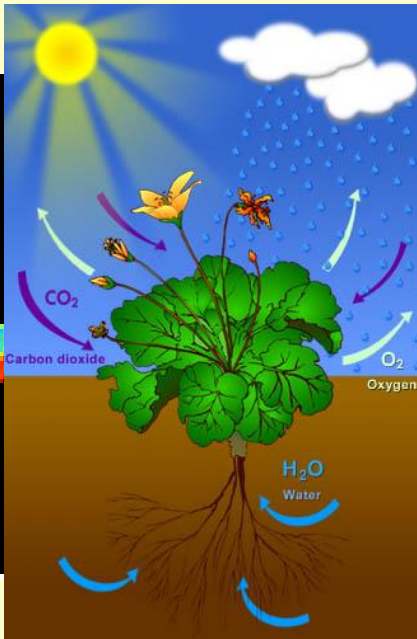
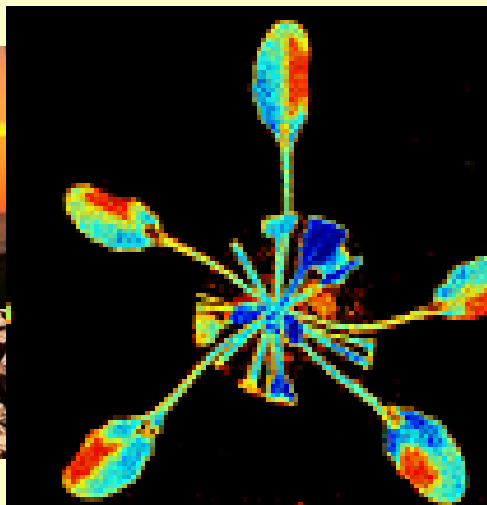




Phenomics

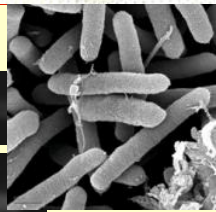
Development and application of high-throughput, multi-dimensional and dynamic phenotyping

- Gene to phenotype modelling -



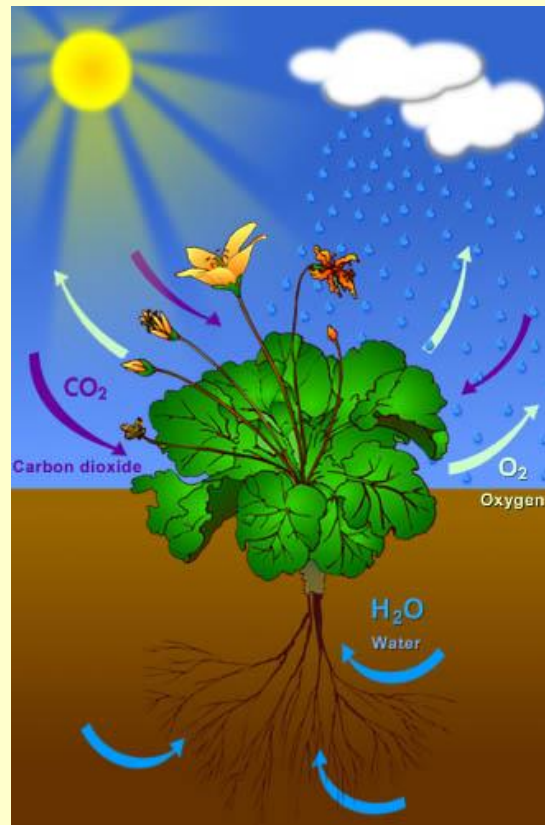
Genexpression

semi-qRT-PCR

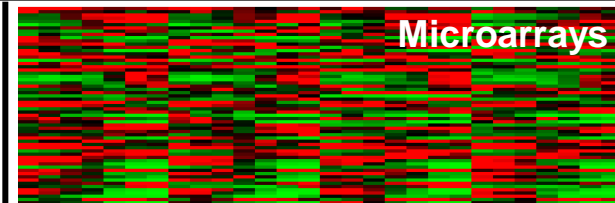
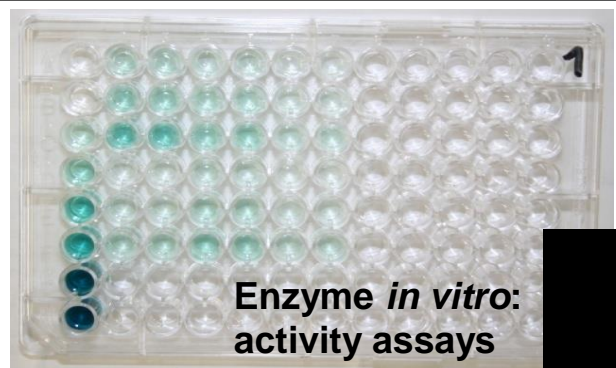


Phenotypic variation

... is produced through a complex web of interactions
between
Genotypes x Environment x Management

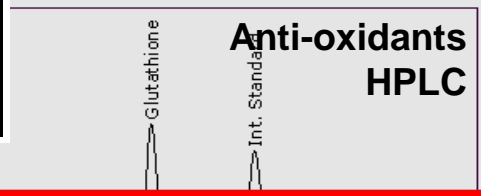
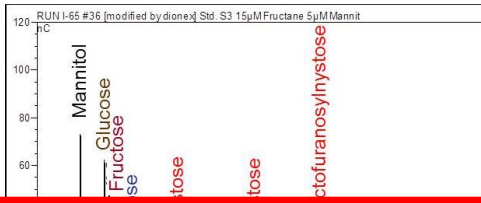
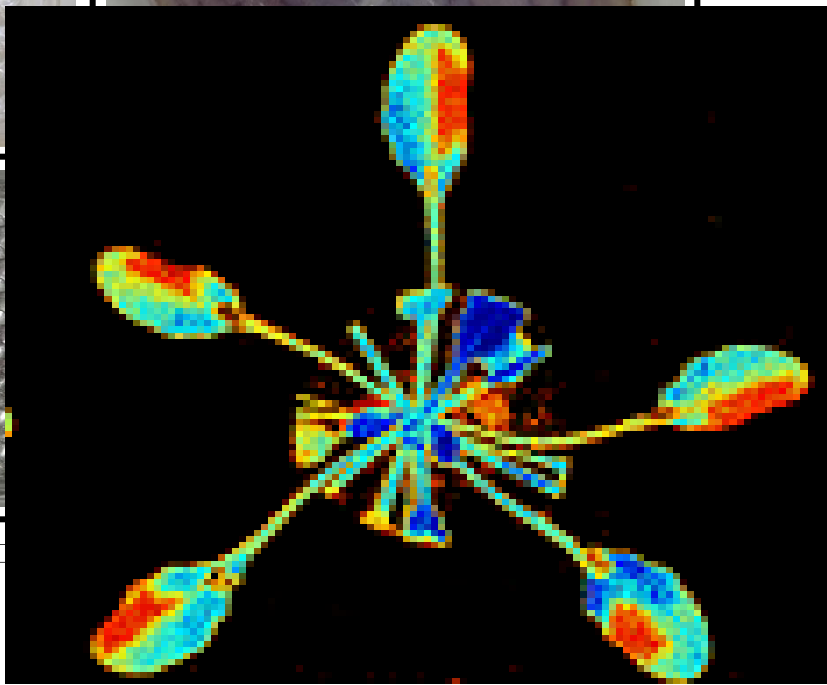
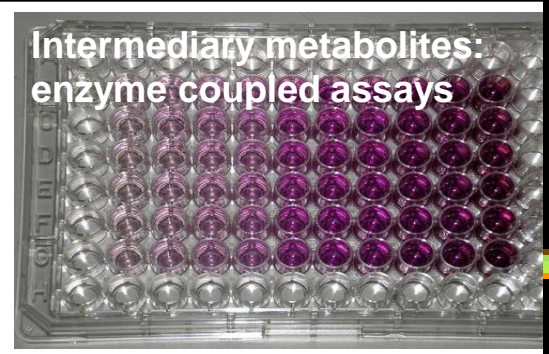


Integration of non-invasive signatures with physiological phenotyping: „Plant physiology meets biophysics“



Genexpression

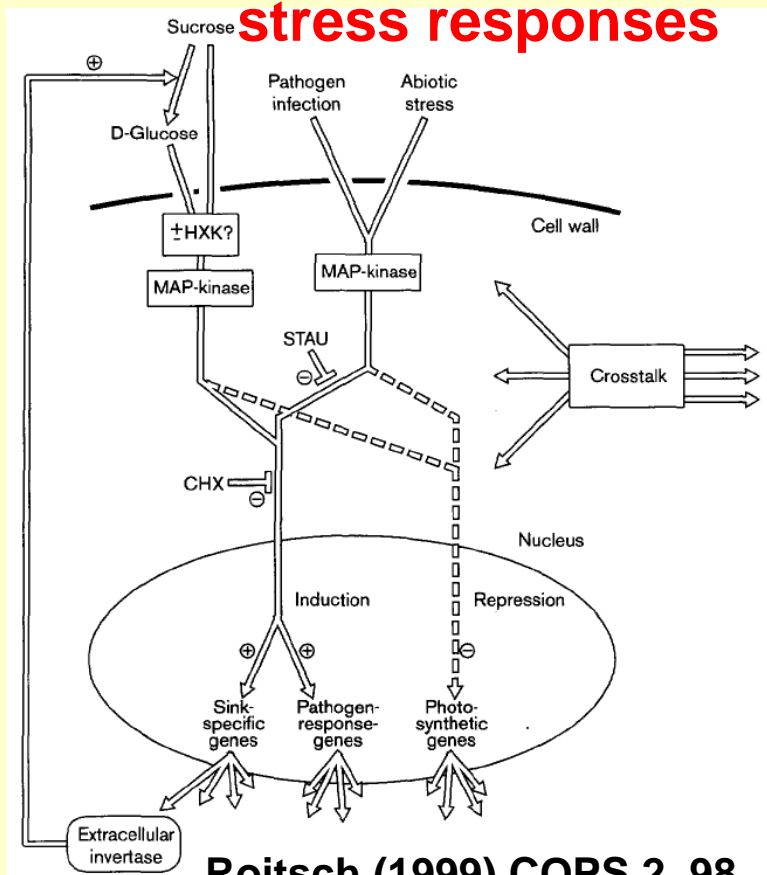
semi-qRT-PCR



Linking hyper-scale, non-invasive imaging to temporal and spatial dynamics of physiological processes and DNA-based markers

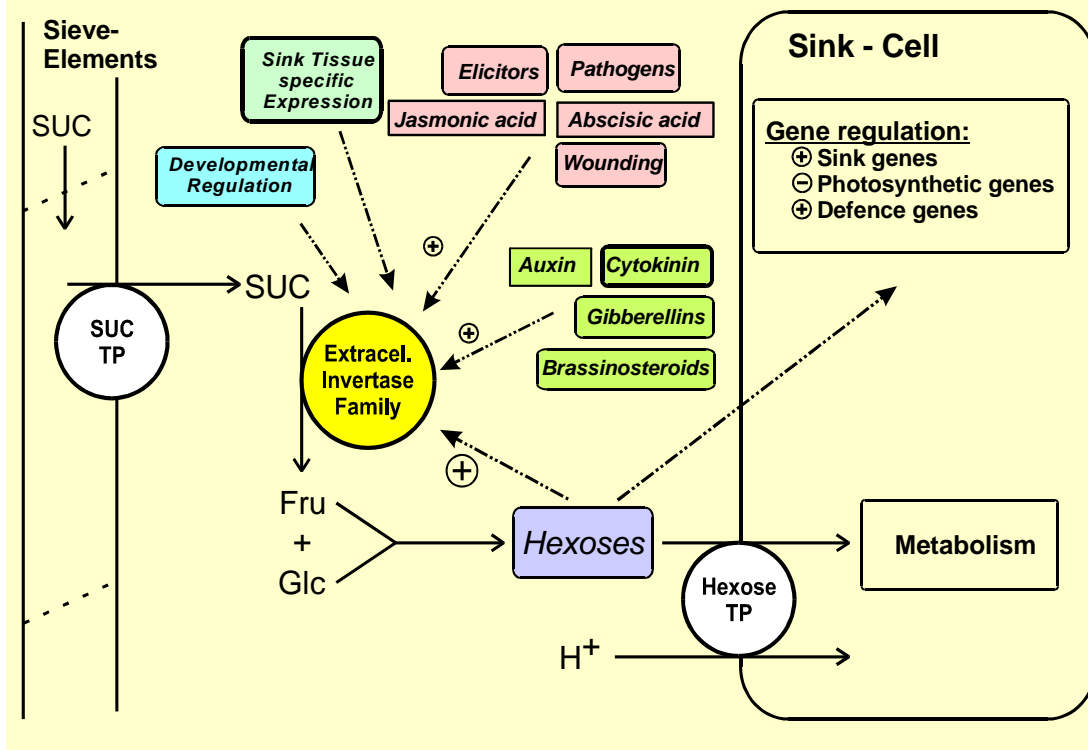
Development & stress responses are linked to plant carbohydrate metabolism

Co-ordinated regulation of source-sink relations & stress responses



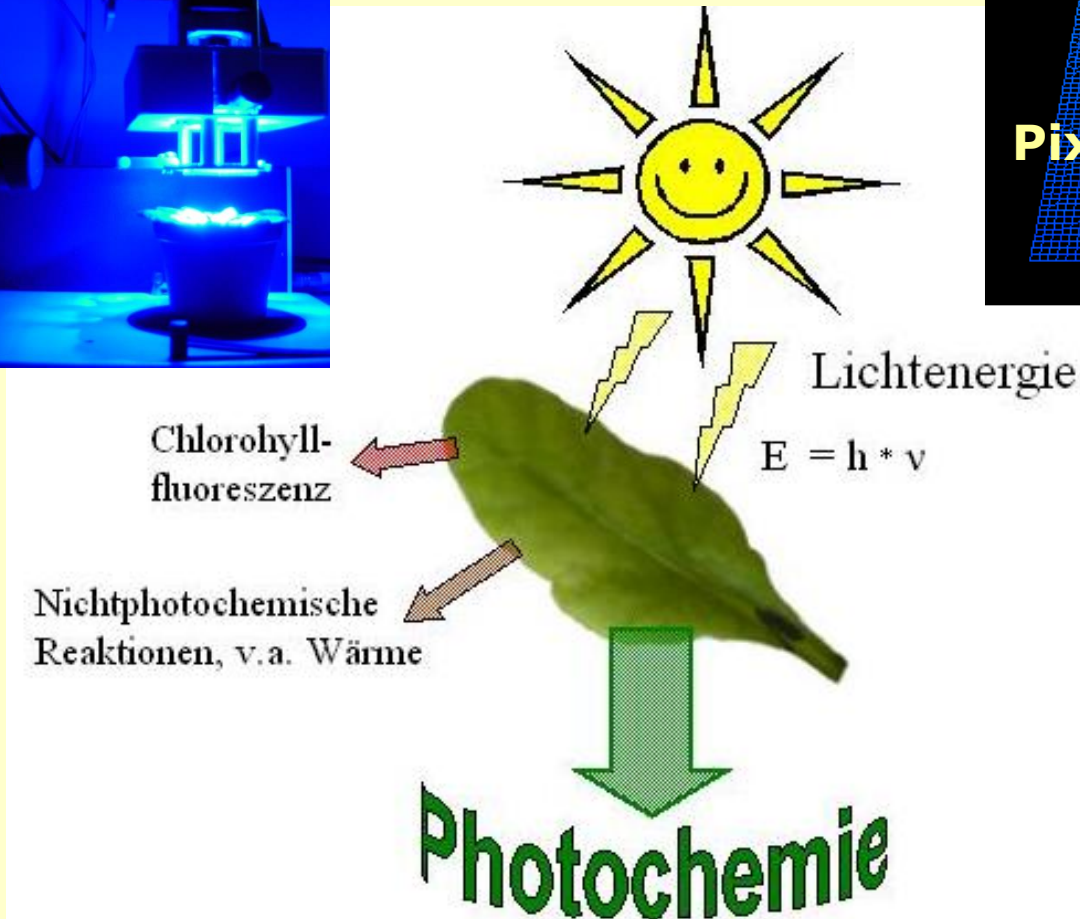
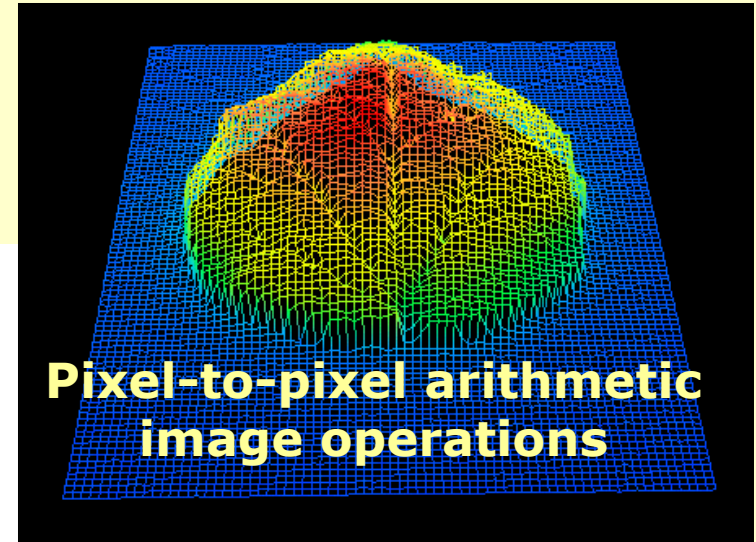
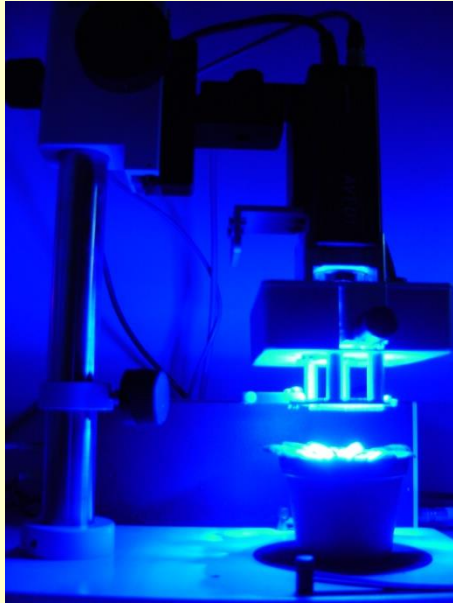
Roitsch (1999) COPS 2, 98
 Berger et al. (2007) JXB 58, 4019
 Albacete et al. (2011) Phyton 50,181

Extracellular Invertase: key metabolic enzyme and stress response protein



Roitsch et al. (1995) PlantPhys 108: 285
 Roitsch et al. (2004) TIPS 9, 607
 Albacete et al. (2011) Phyton 50,181

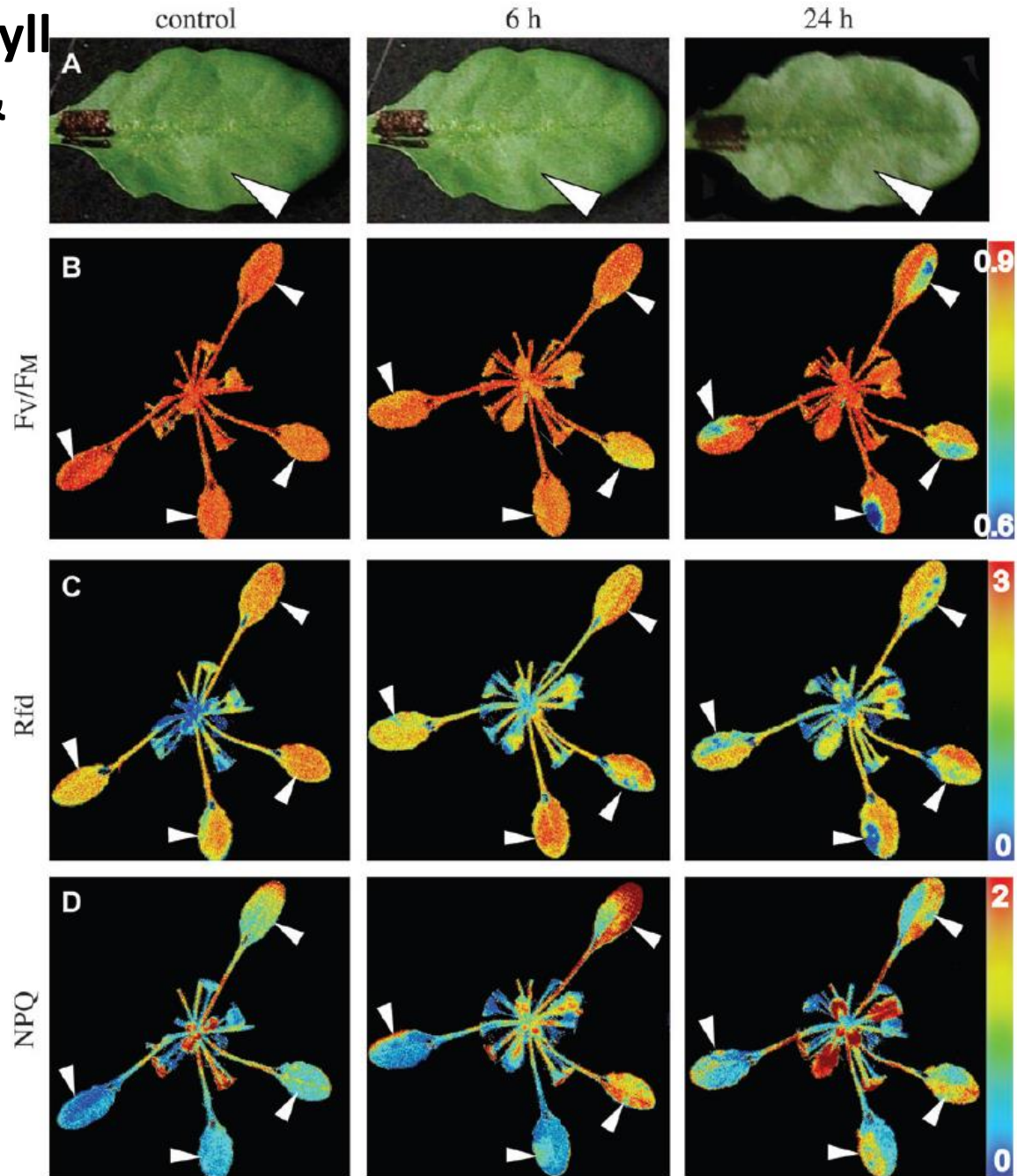
Chlorophyll Fluorescence Imaging



Advantages:

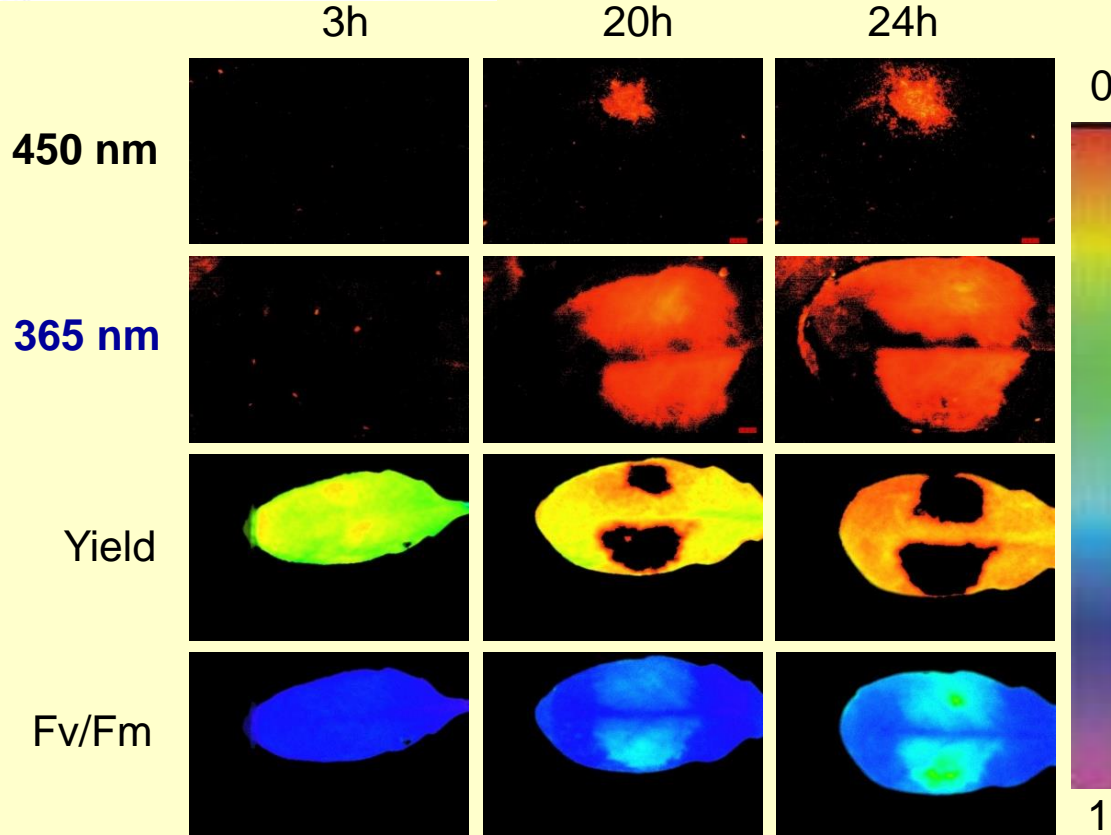
- Non invasive
- Very high sensitivity
- Spatial resolution

Combination of chlorophyll fluorescence imaging & statistical analysis: Combinatorial imaging



Development of multifluorescence imaging

Discrimination between chlorophyll, phenolic and GFP fluorescence



Simultaneous infiltration of WT strain and GFP labelled strain of *P. syringae*

PAM/GFP- measuring head

UV-light measuring head

PAM/GFP- measuring head

Enzyme activity signatures

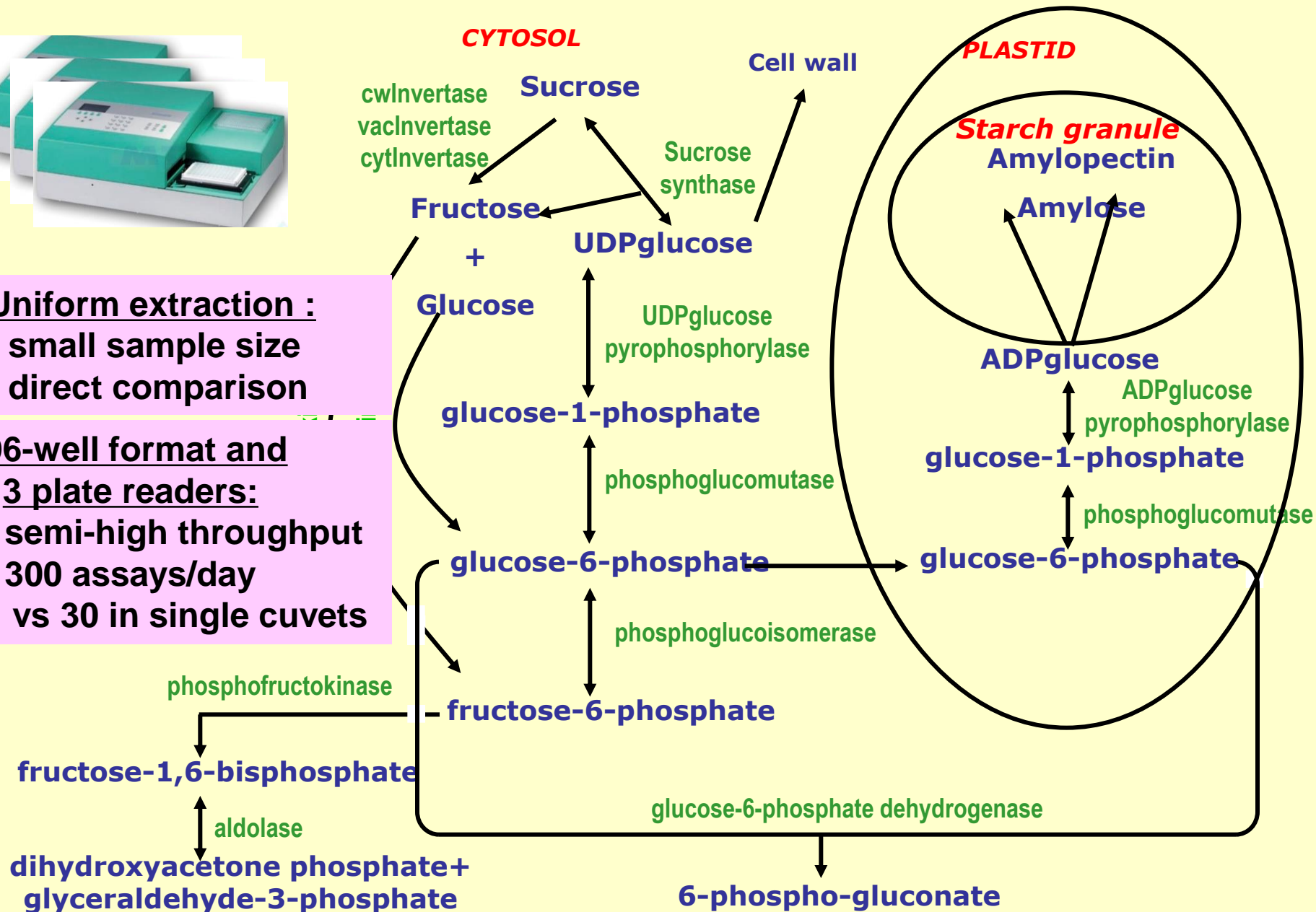


Uniform extraction :

- small sample size
- direct comparison

96-well format and 3 plate readers:

- semi-high throughput
- 300 assays/day vs 30 in single cuvetts



Characterisation of sugar beet development

Times course of enzyme activity signatures

Parameter [min/max]	No. 1	No. 2	No. 3	No. 4
Frischgewicht (0,01 – 25 g)	▲ ▲ ▲ ▲ ▲ ▲	▲ ▲ ▲ ▲ ▲ ▲	▲ ▲ ▲ ▲ ▲ ▲	▲ ▲ ▲ ▲ ▲ ▲
Suc-Gehalt (10 – 60.000 µg/gFW)	▲ ▲ x ▲ ▲ ?	▲ ▲ x ▲ ▲ ?	▲ ▲ x ▲ ▲ ?	▲ ▲ x ▲ ▲ ?
Sucrol. Aktiv. ●	▼ ▼ ▼ ▼ ▼ ▼	▼ ▼ ▼ ▼ ▼ ▼	▼ ▼ ▼ ▼ ▼ ▼	▼ ▼ ▼ ▼ ▼ ▼
vacInv ●	▼ ▼ ▼ ▼ ▼ ▼	▼ ▼ ▼ ▼ ▼ ▼	▼ ▼ ▼ ▼ ▼ ▼	▼ ▼ ▼ ▼ ▼ ▼
cytInv ●	▼ ▼ ▼ ▼ ▼ ▼	▼ ▼ ▼ ▼ ▼ ▼	▼ ▼ ▼ ▼ ▼ ▼	▼ ▼ ▼ ▼ ▼ ▼
cwlInv ●	▼ ▼ ▼ ▼ ▼ ▼	▼ ▼ ▼ ▼ ▼ ▼	▼ ▼ ▼ ▼ ▼ ▼	▼ ▼ ▼ ▼ ▼ ▼
SuSy (nd – 2 nkat*mg ⁻¹ Prot) ●	▲ ▲ ▲ ▲ ▲ ▲	▲ ▲ – ▲ ▲ ▲	▲ – – ▲ ▲ ▲	▲ – – ▲ ▲ ▲
PFK (nd – 1 nkat*mg ⁻¹ Prot) ●	▲ ▲ – ▲ ▲ ▲	▲ ▲ – ▲ ▲ ▲	▲ ▲ ▲ ▲ ▲ ▲	▲ ▲ ▲ ▲ ▲ ▲
UGPase (nd – 6 nkat*mg ⁻¹ Prot) ●	▲ ▲ ▲ ▲ ▲ ▲	▲ ▲ ▲ ▲ ▲ ▲	▲ ▲ ▲ ▲ ▲ ▲	▲ ▲ ▲ ▲ ▲ ▲
PGI (nd – 20 nkat*mg ⁻¹ Prot) ●	▲ ▲ ▲ ▲ ▲ ▲	▲ ▲ ▲ ▲ ▲ ▲	▲ ▲ ▲ ▲ ▲ ▲	▲ ▲ ▲ ▲ ▲ ▲
PGM (nd – 20 nkat*mg ⁻¹ Prot) ●	▲ ▲ ▲ ▲ ▲ ▲	▲ ▲ ▲ ▲ ▲ ▲	▲ ▲ ▲ ▲ ▲ ▲	▲ ▲ ▲ ▲ ▲ ▲
G6PDH (nd – 3 nkat*mg ⁻¹ Prot) ●	▲ ▲ ▲ ▲ ▲ ▲	▲ ▲ ▲ ▲ ▲ ▲	▲ ▲ ▲ ▲ ▲ ▲	▲ ▲ ▲ ▲ ▲ ▲
Aldolase (< 0,1 – 2 nkat*mg ⁻¹ Prot) ●	∩ – ∩ ∩ ∩ ∩	∩ – ∩ ∩ ∩ ∩	∩ ∩ ∩ ∩ ∩ ∩	∩ ∩ ∩ ∩ ∩ ∩

▲ = Increase over time (90 days); ▼ = decrease over time; ∩ = transient;
 – = no change; x = n. d.; ? = no data yet

Instrumental analysis: Hormone levels

1. Cytokinins

a) biological active

- *trans*-Zeatin (tZ)
- *trans*-Zeatin-Ribosid (tZR)
- Dihydrozeatin (DHZ)
- Dihydrozeatin-Ribosid (DZR)
- Isopentenyl-Adenin (iP)
- *cis*-Zeatin (cZ)

b) conjugates:

- Isopentenyl-Adenin-7-Glucoside (IP7G)
- *trans*-Zeatin-O-Glucosid (ZOG)
- *trans*-Zeatin-O-Glucosid-Ribosid (ZOGR)
- *trans*-Zeatin-7-Glucoside (tZ7G)
- *trans*-Zeatin-9-Glucoside (tZ7G)

2) other hormones/ signal molecules

- Indolacetic Acid (IAA)
- Abscisic Acid (ABA)
- Jasmonic Acid (JA)
- Salicylic Acid (SA)

3) To be established: Gibberellins

- G A1
- G A3
- G A4
- G A5
- G A6
- G A8

UHPLC/Ms-Ms



Determination of spatial and temporal dynamics of plant responses during development within environmental constraints and interactions

