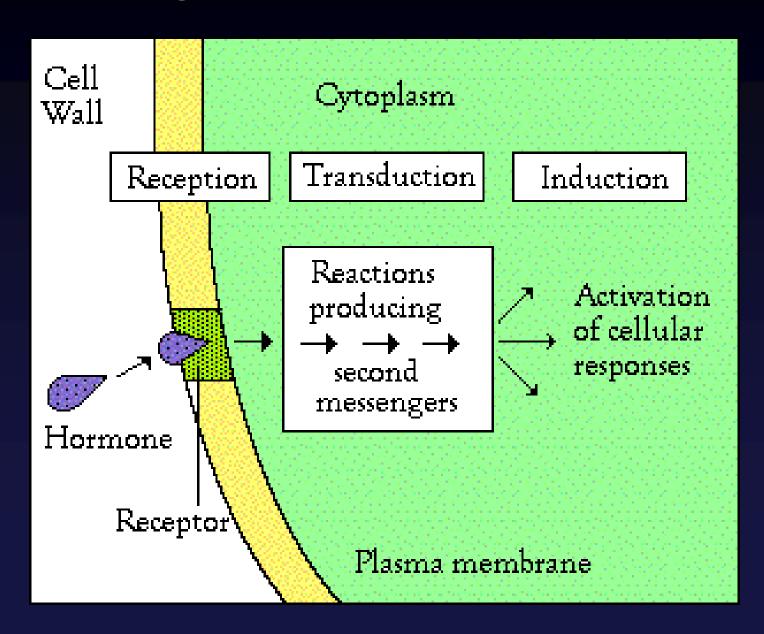
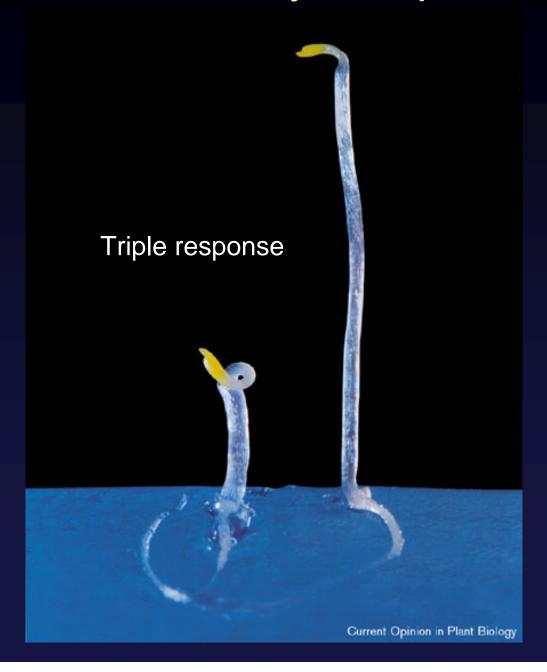
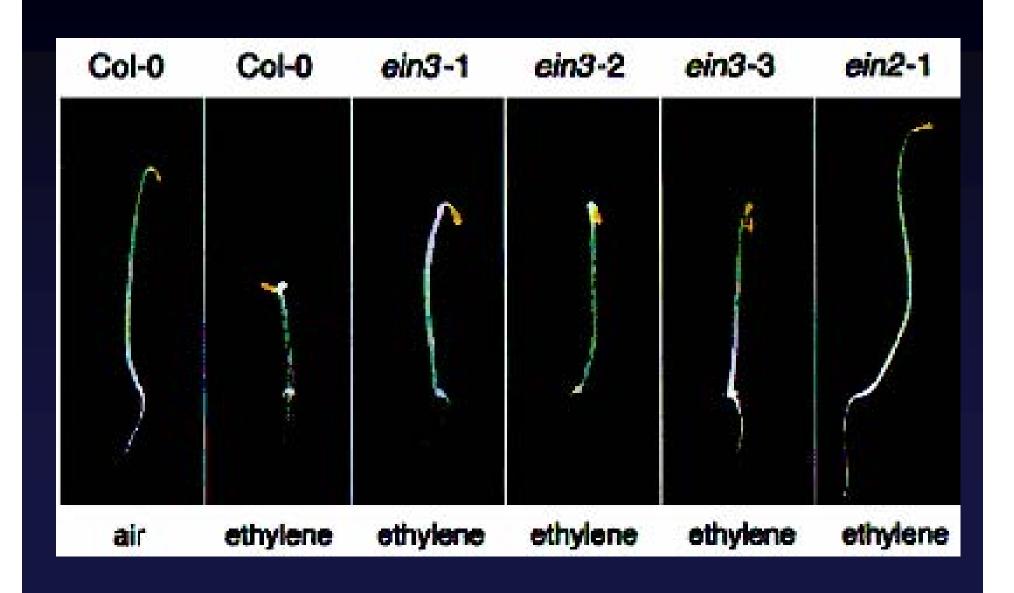
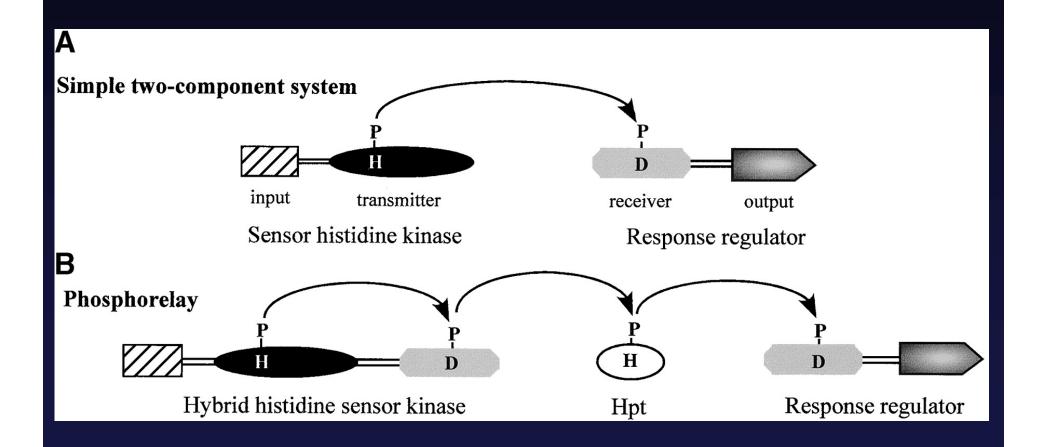
Signal Transduction

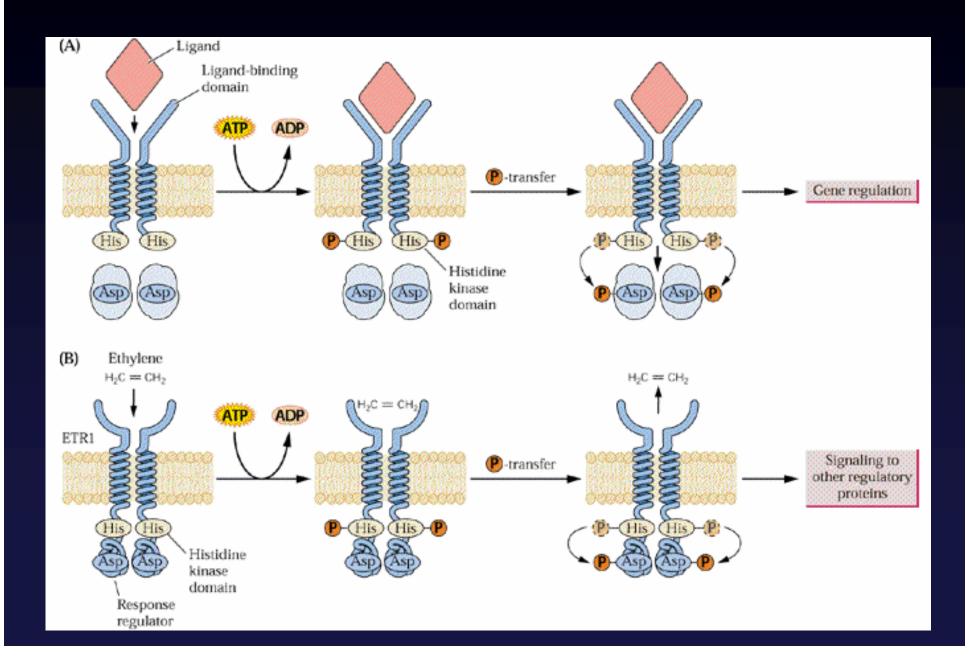


Mutant screens for ethylene pathway genes

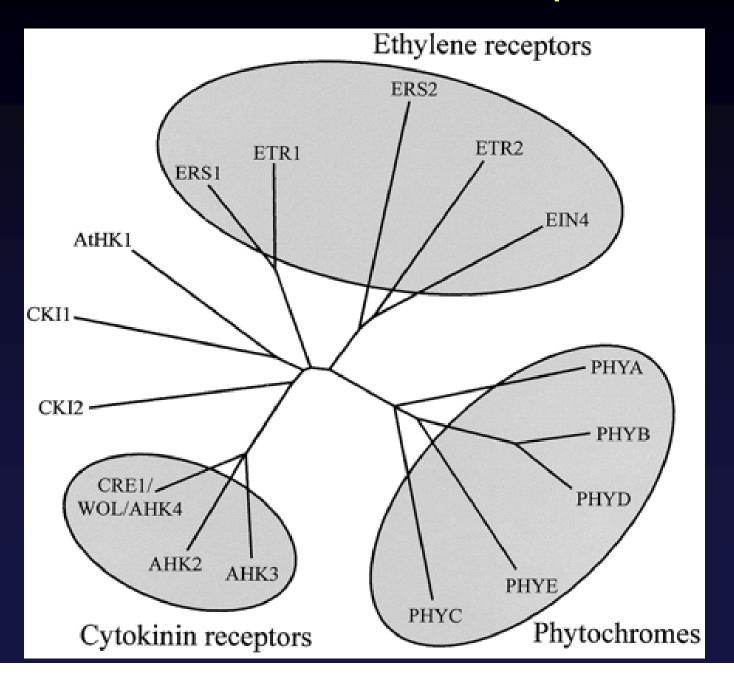




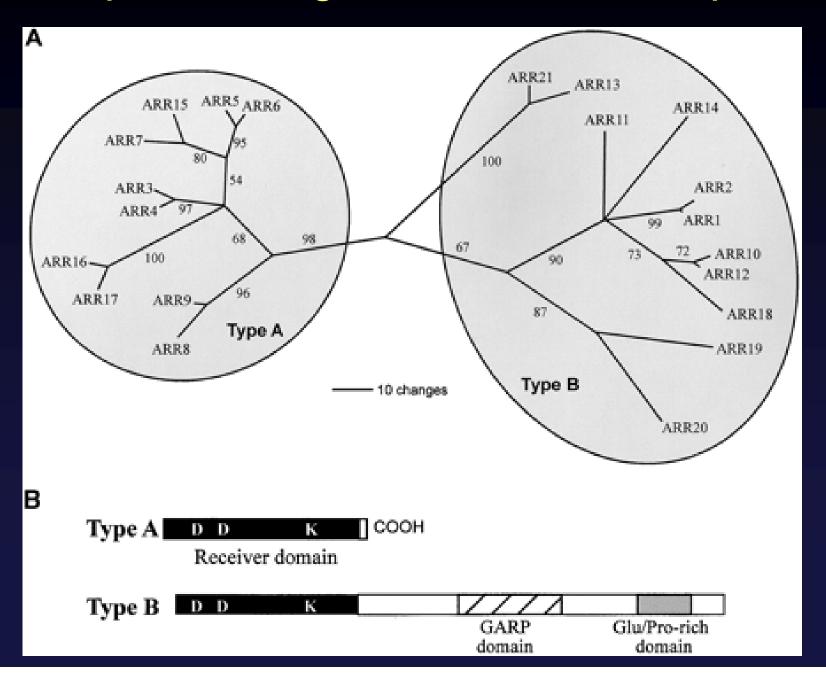


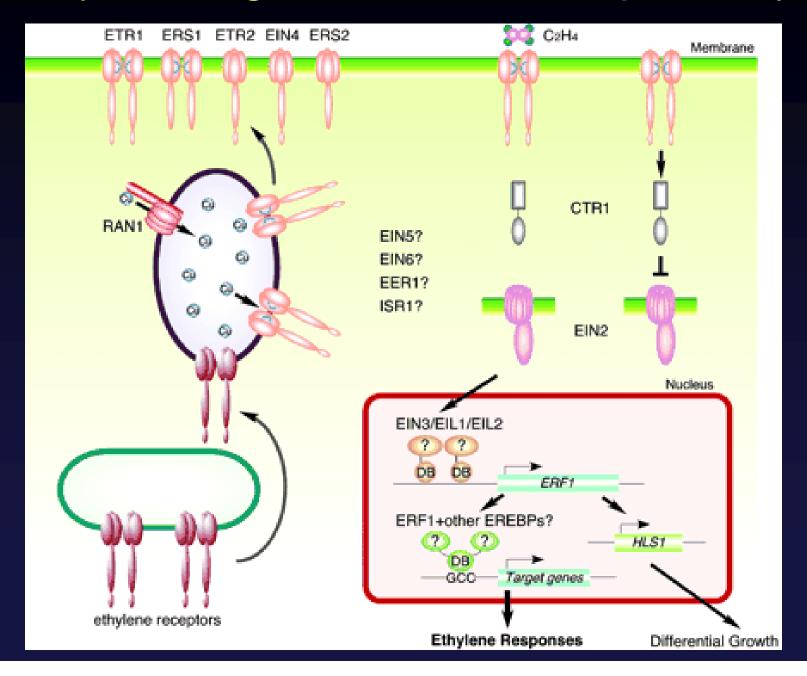


His-kinases in Arabidopsis

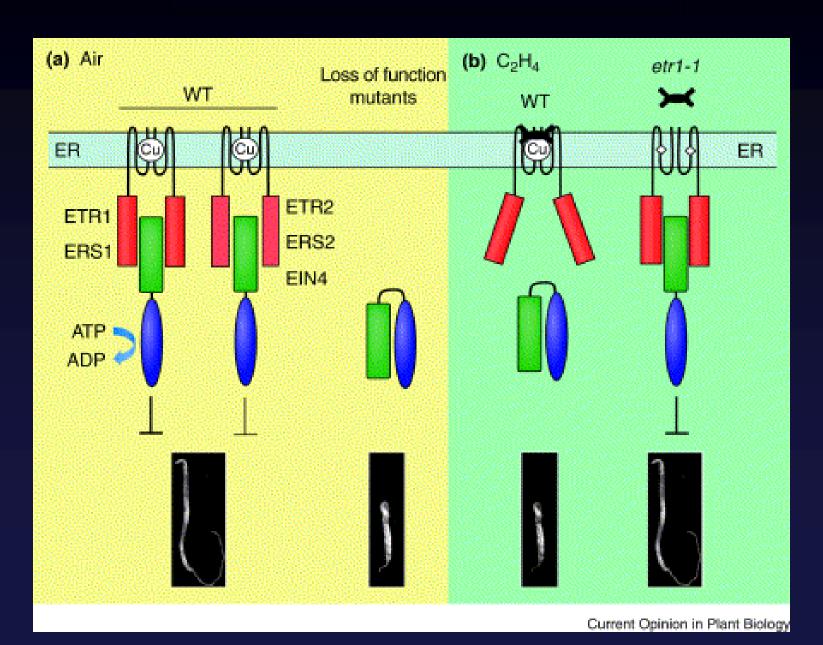


Response Regulators in Arabidopsis





Genetic interactions



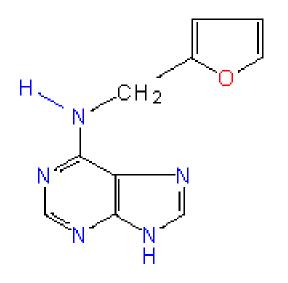
CYTOKININ – what is important?

Synthesis – IPT genes

Degradation – CK-oxidase

Signal transduction – forward genetics activation tagging CK regulated genes

Cytokinins



Kinetin:

6 - (2 - Furfuryl -

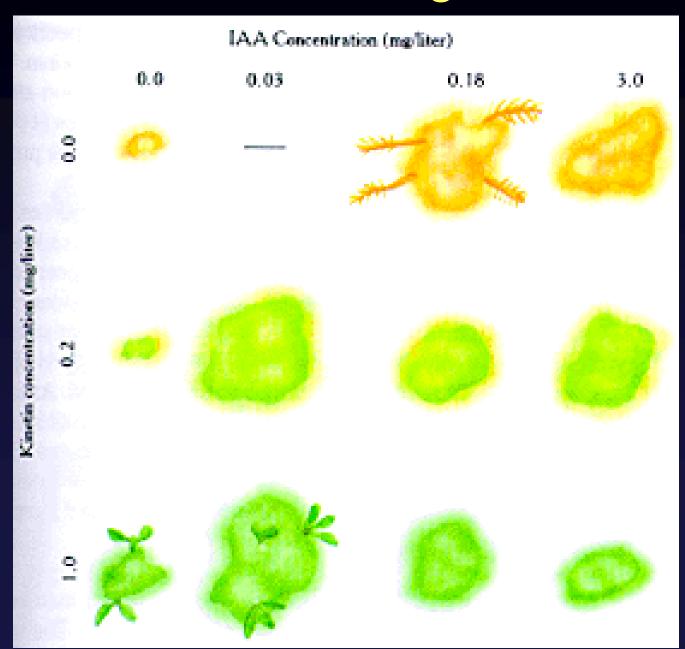
7 - Aminopurin)

Cytokinin (Grundstruktur)

Zeatin

$$CH_2$$
OH $-CH_2$ — CH_2 — CH_2 — CH_3
Dihydrozeatin CH_3

Effect of CK on regeneration

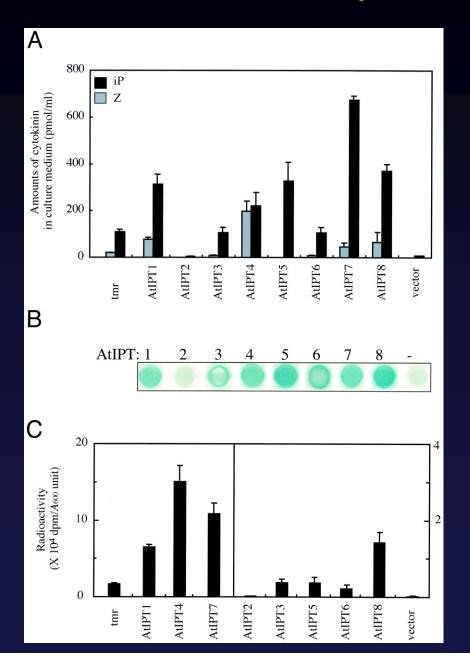


Manipulating of CK levels by overexpression of bacterial *IPT*

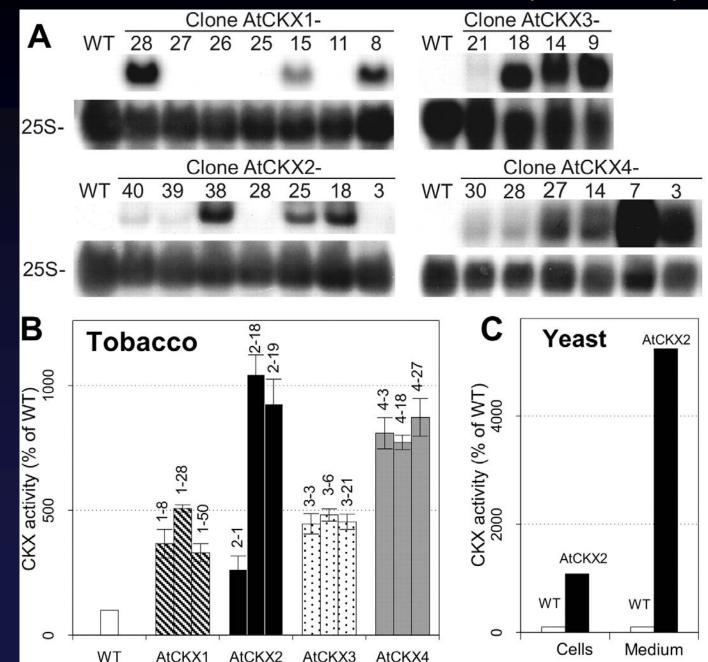


Region a MTELNFHLLPIISDRFTTTTTTSPSFSSHSSSSSSLLSFTKERRKHOPLVSSIRMEQSRSRURKD MMMLNPSNGGIEGEKMKKKA MIMKISMAMCKOPEPPSPTLDFPPARFGPNMLTLN-PY MKCS. MKPCMTALRQVIQPLSLNFQGNMVDVP-FFRRR MQQLMTLLSPPLHHSSLLPTVTTKFGSPRLVTTCHGHAGRKEIT, MKFSISSLKQVQPILCFKNKLSKVNVNSFLHPR MQNLTSTVGFMNPITSPHLBLPPRSVVPMTTVCMCQSY AtIPT8 VVSKFLLDDAAEDTEECCADVASVVDQDMVVESVFGRDDLSHGYELLKELDPVAANRIHPNNHRKINGY AIRWOALRAMEEAIDA NEPKDOKLRIMEEAIDA -FLANKOBEELLSKULEEARE -DRWDPMRKEAYEKAVRAIKE OF -NYPAETTERLEETAIEKU OF -NYPAETTERLEETAIEKU OF -NYPAETTERLEETAIKKO EK #KAVHOKSSS - EMANGEN ELQULBENYKLYKRILMOS LIS - SEESGAANVIKANSE IKCPERTEEGORDPTSO EKERGAANVIKANSE IKCPERTEEGORDPTSO EKERLINAAKSOKANSE IKCPERTEEN ELDER EVITAS FERRITAAKSOKANSE IKKERIS VERVENSE OPENYKYTYNS FERRITAAKSOKANSE IKKERIS VERVENSE OPENYKYTES FERRITAAKSOKANSE IKKERIS VERVENSE OPENYKYTES VERVENSE FERRITAAKSOKANSE IKKERIS VERVENSE VER AtIPT2 AtIPT3 AtIPT4 311 LKRFLSLN 311 LLPEISAVPPLPAAVAAISR 312 CLAASYGGGSGSRAHNMI В Prokaryotic R. prowazekii M. leprae S. coelicolor tRNA-IPT Eukaryotic C. elegans tRNA-IPT AtIPT2 PCC6803 AtIPT Bacterial pTi-SAKURA A. tumefaciens IPT

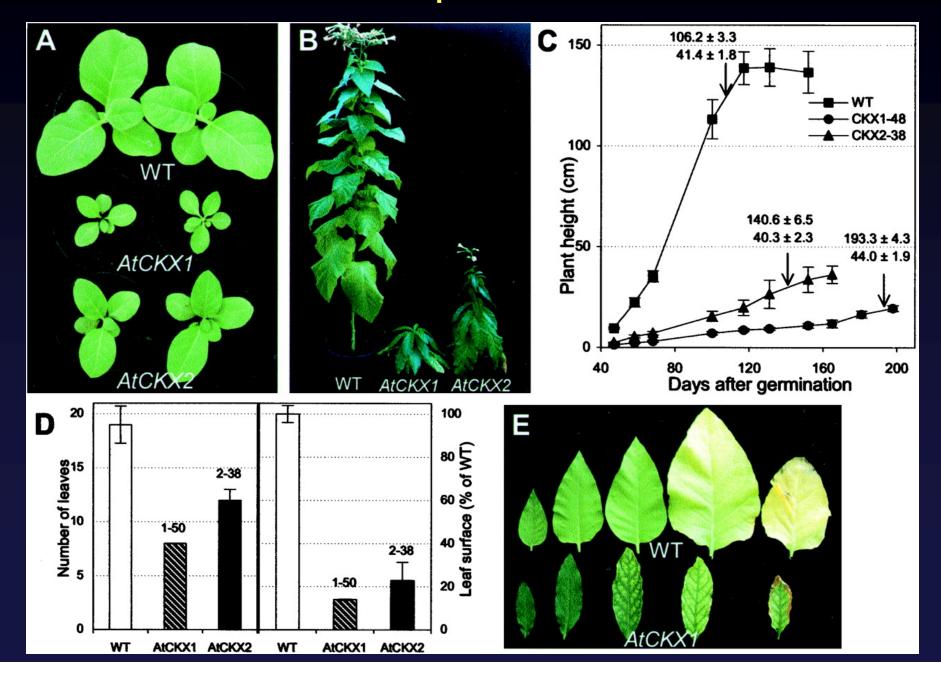
Isolation of Arabidopsis *IPT*s



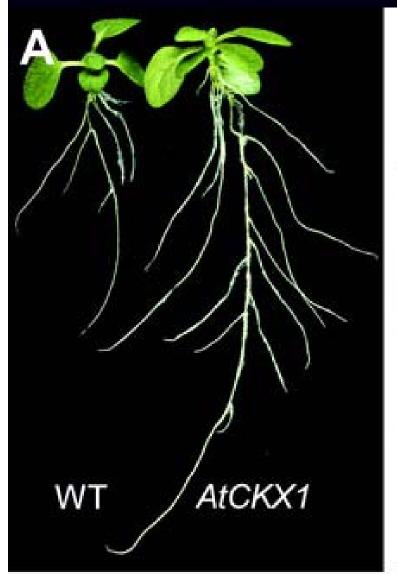
Isolation of CK-oxidase (AtCKX)



AtCKXs overexpression in tobacco

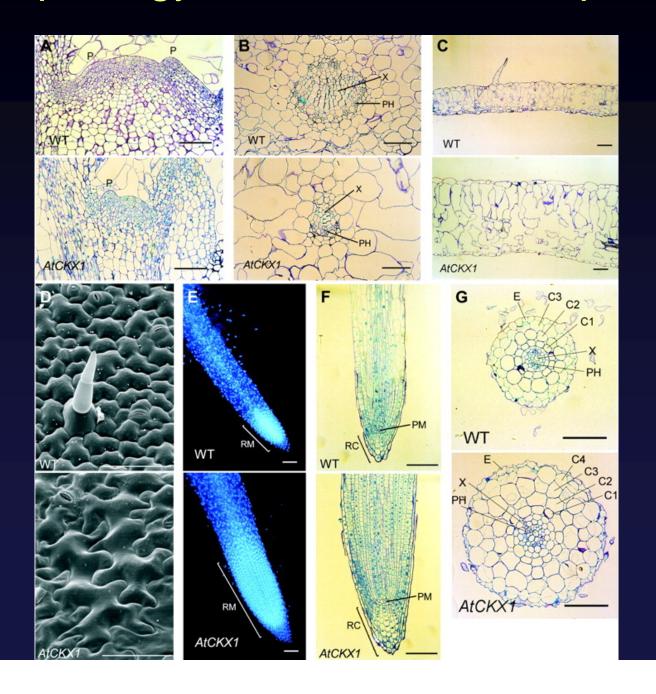


Effect of AtCKX on tobacco root

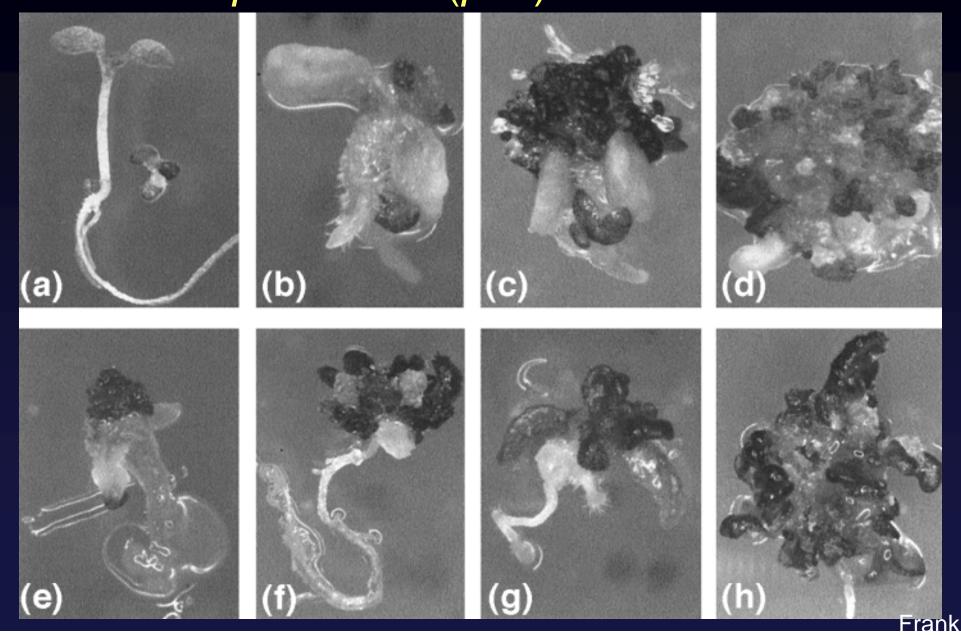




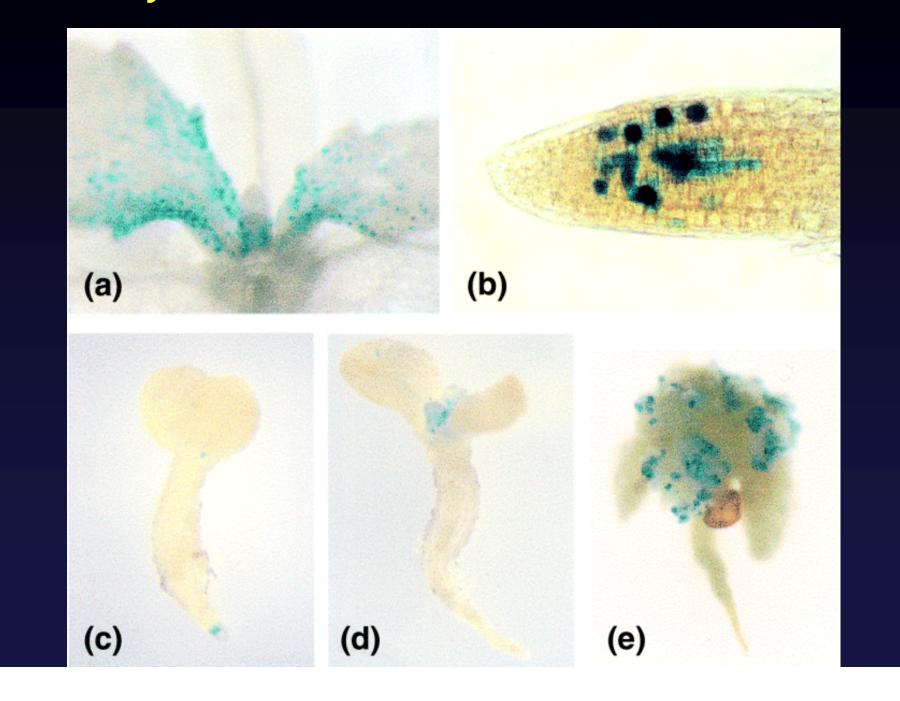
Morphology of AtCKX tobacco plants



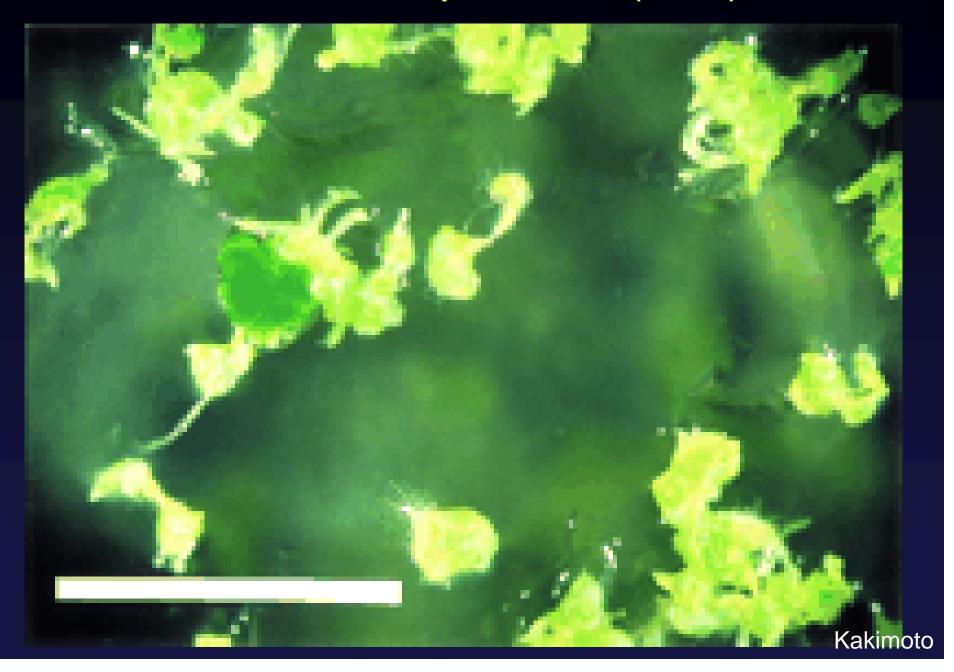
Tumor shoot development (*tsd*) and *pasticcino* (*pas*) mutants



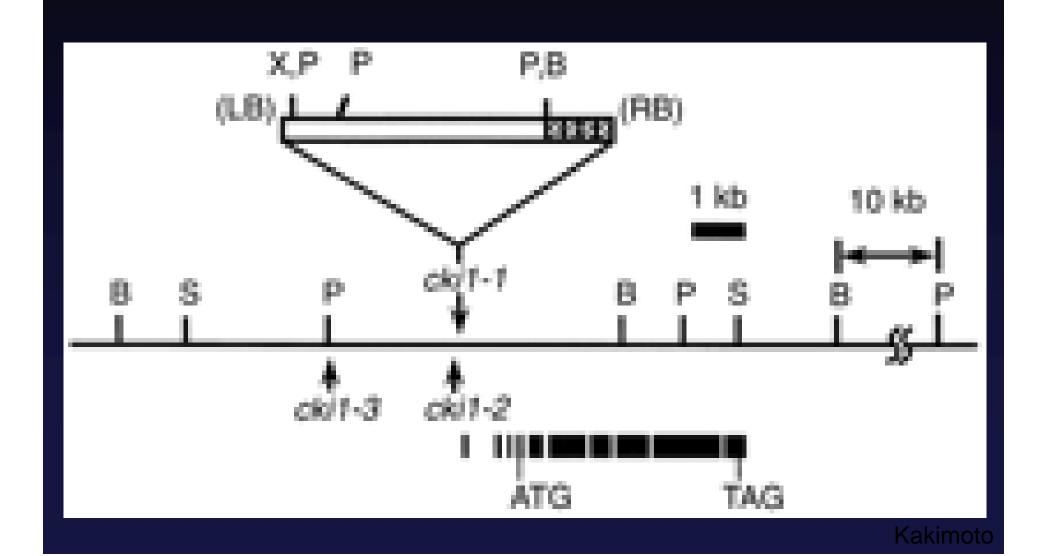
Use of CyclinB::GUS for cell division monitoring



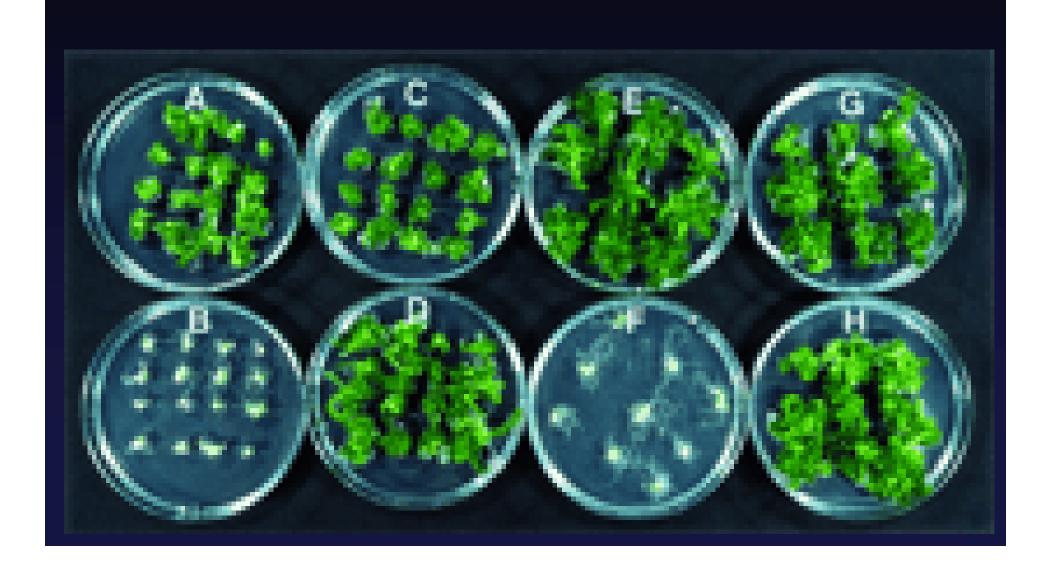
Isolation of CK independent (cki1) mutant

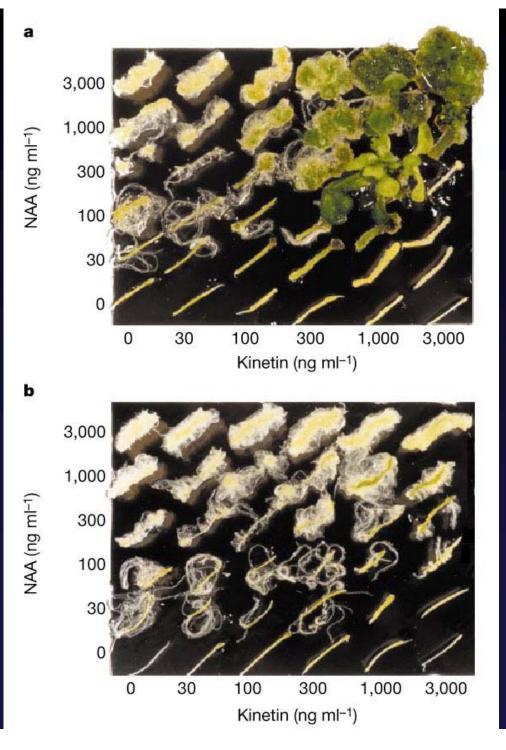


Identification of CKI1 gene



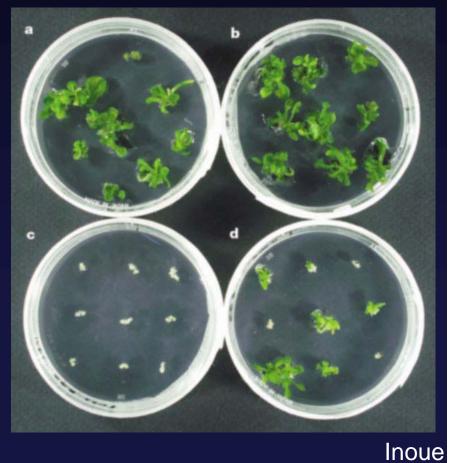
Verification - 35S::CKI1 transgene



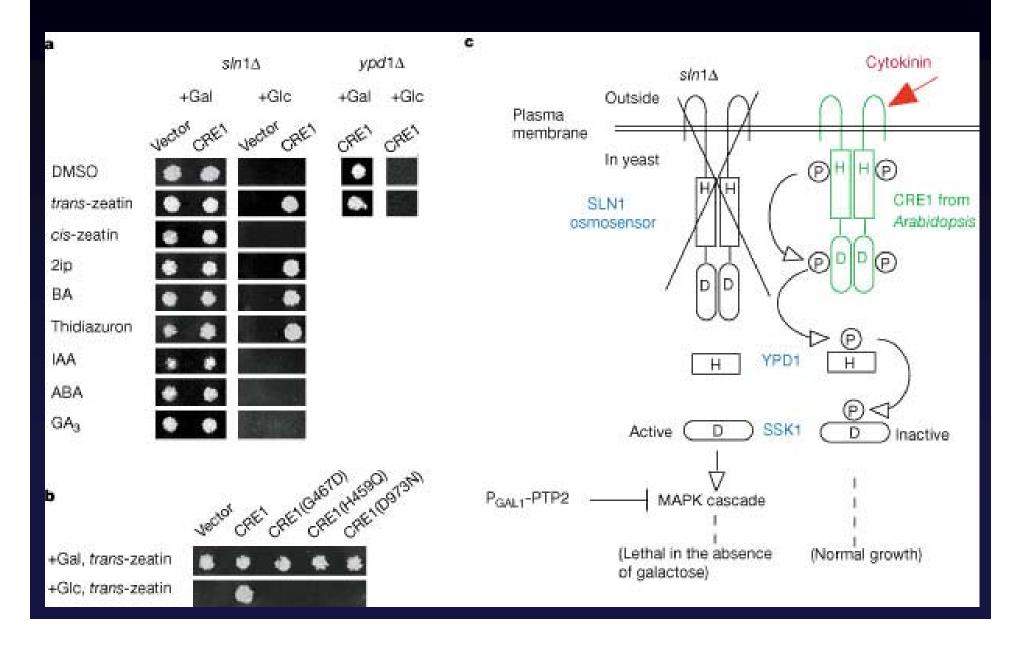


Next strike

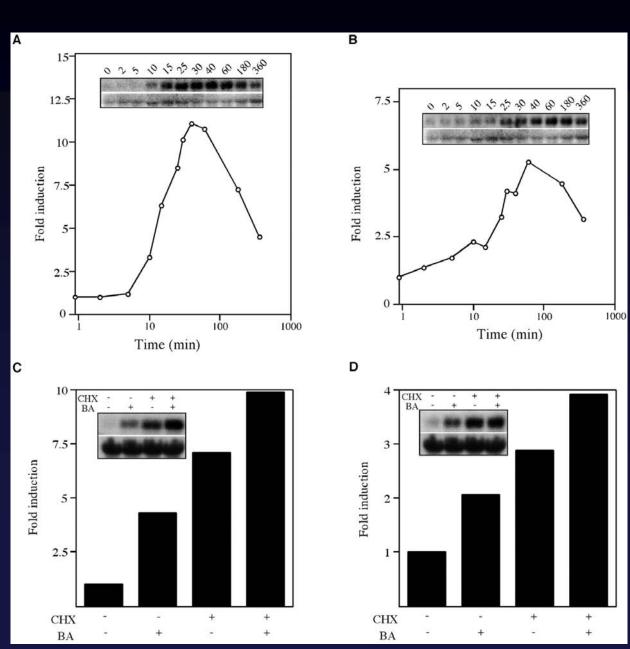
- CK response mutant (cre1)

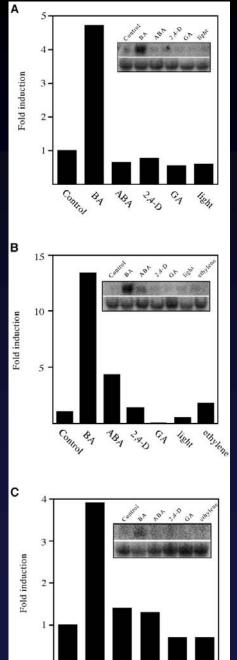


Piece of genius - complementation



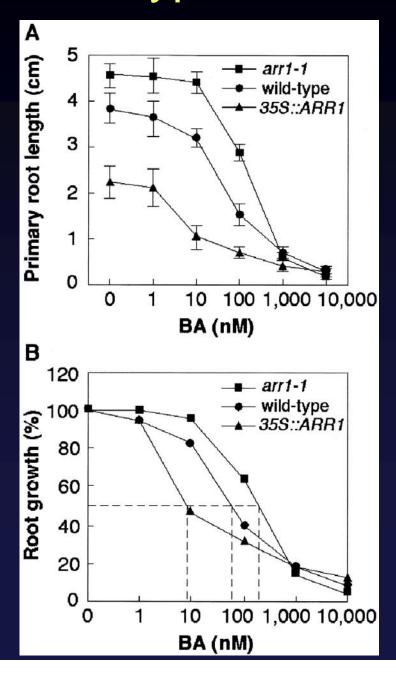
CK responsive genes - ARRs

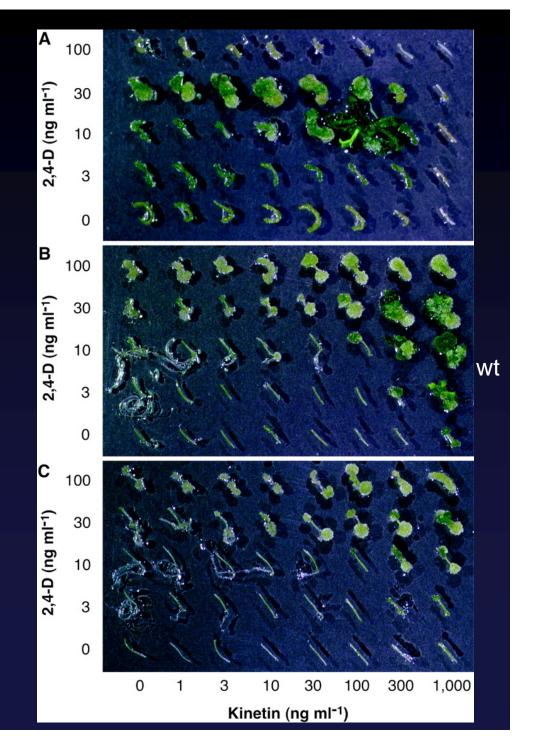




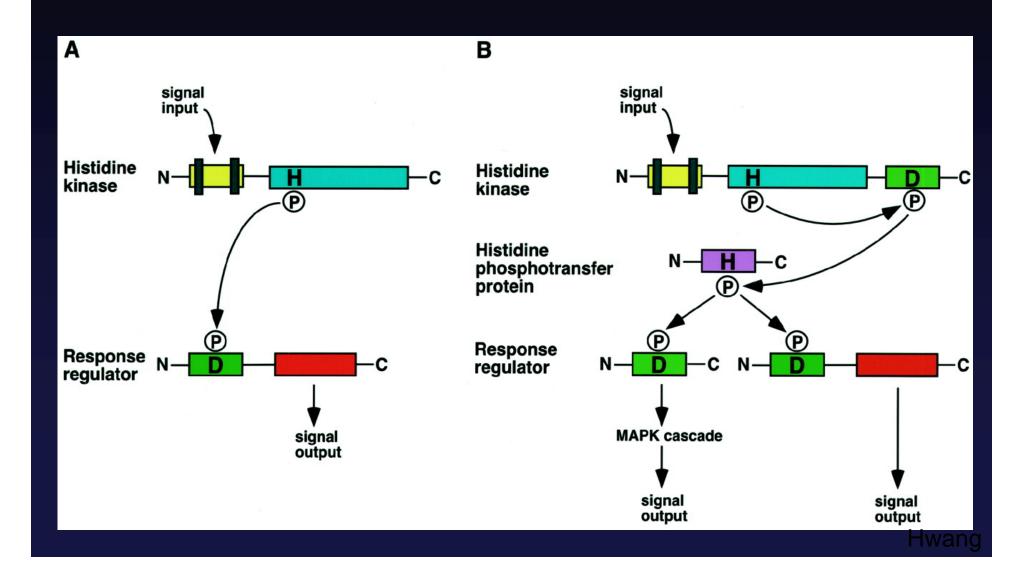
ndstatter

Phenotypes of arrs

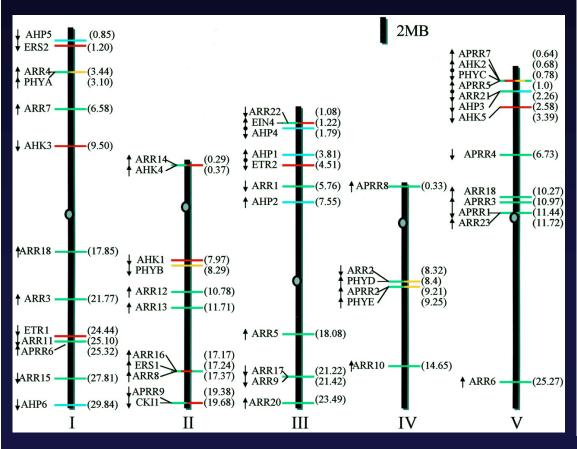


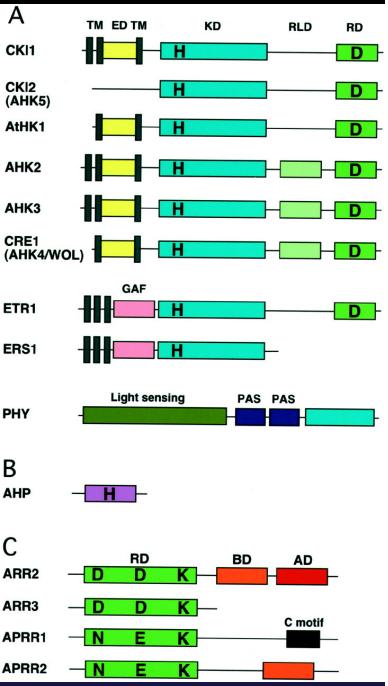


His kinase transduction pathway



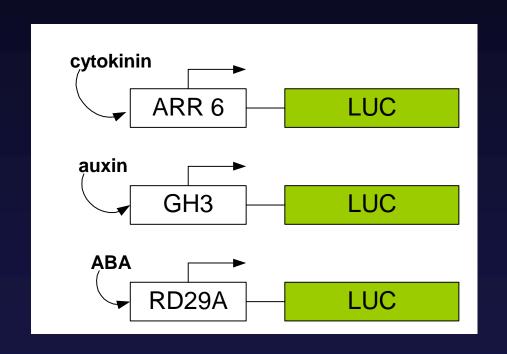
His kinase pathway components in Arabidopsis

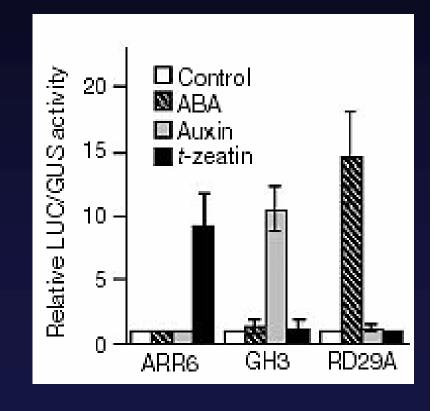




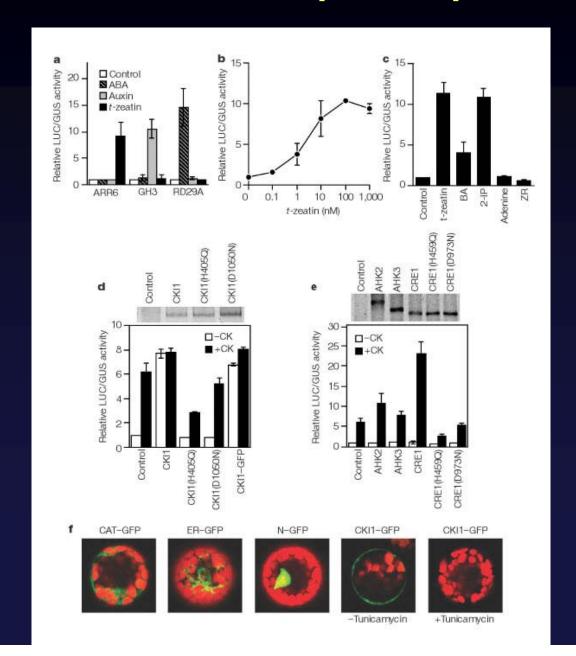
Games with protoplasts Poportors for different pathwo



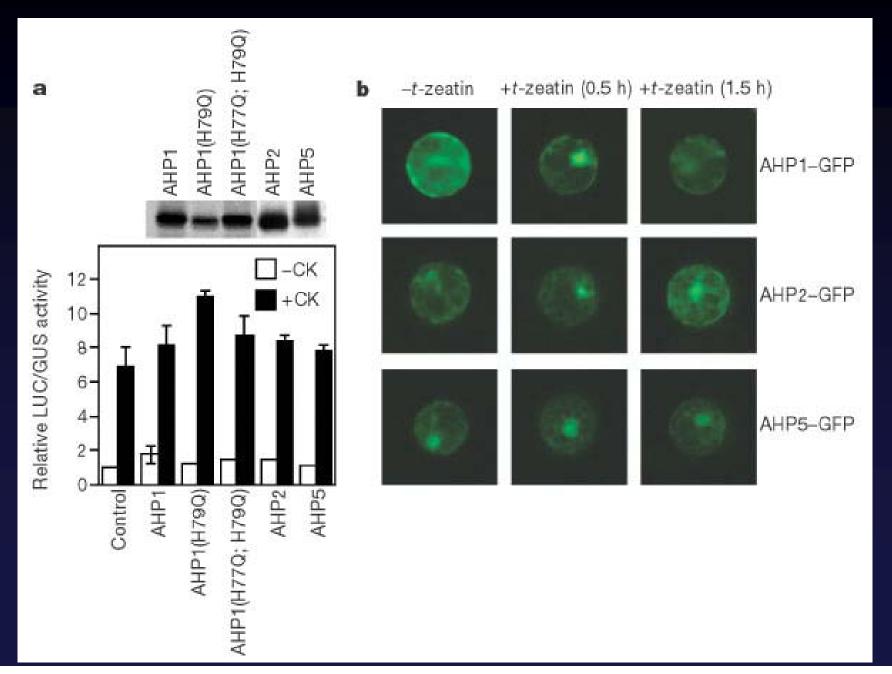




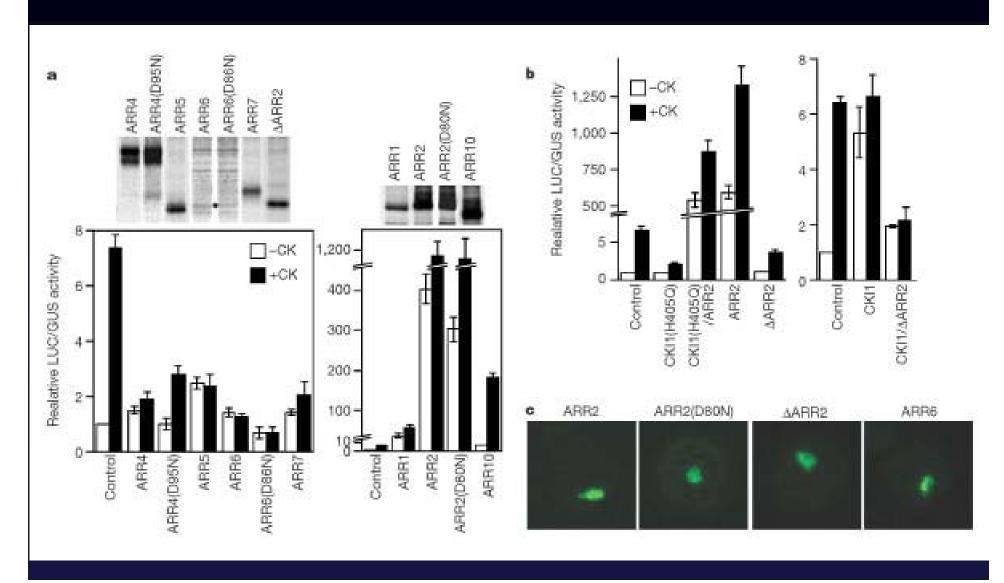
Games with protoplasts



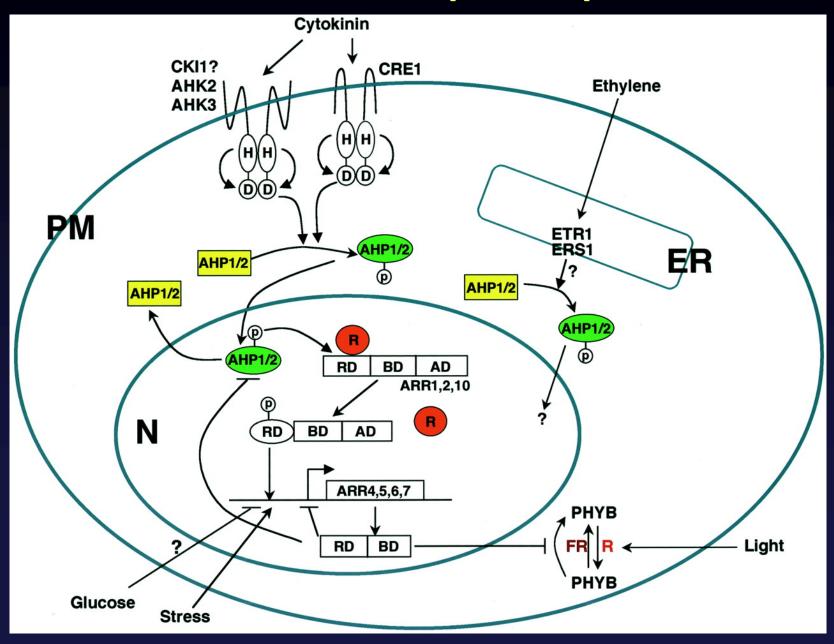
AHPs – shuttle to nucleus



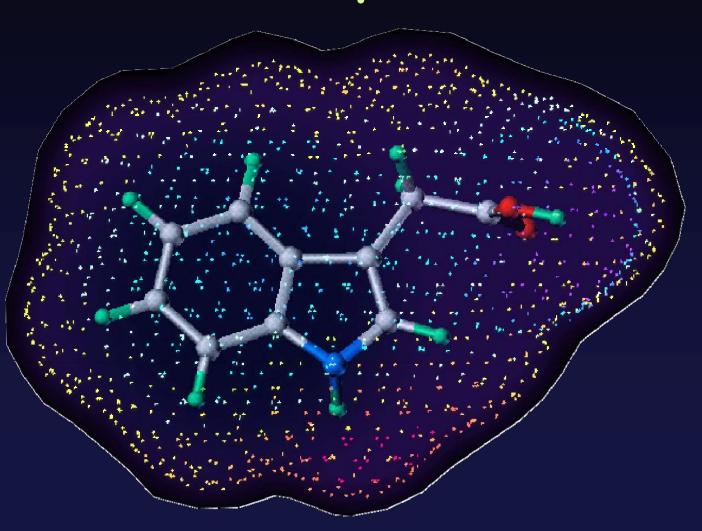
Opposite effects of two classes of ARRs on CK signalling



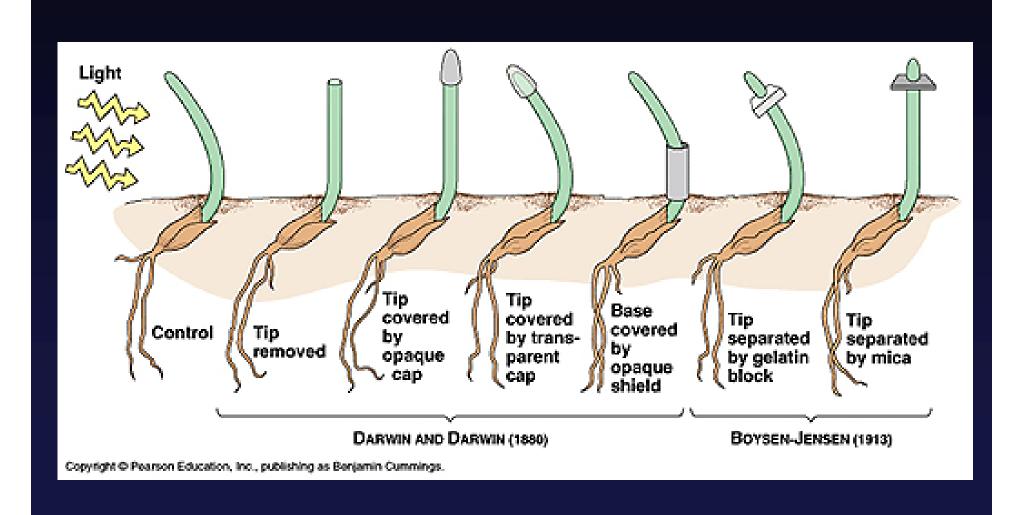
Games with protoplasts



Auxin Signaling and Transport

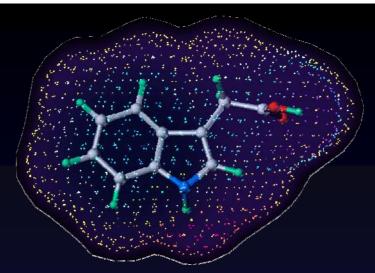


Discovery of the First Plant Signaling Molecule – Auxin and its Transport



AUXIN

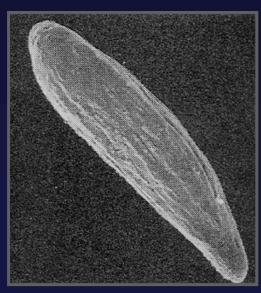
mediates



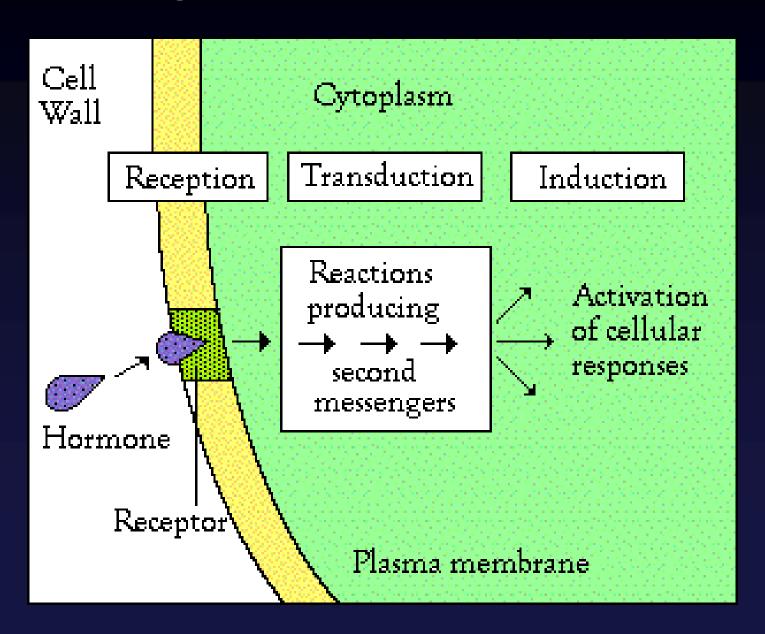
- Embryo development
- Organ initiation and positioning
- Vascular tissue differentiation
- Shoot and root elongation
- Growth responses to light and gravity
- Apical hook formation

embryos





Signal Transduction



Biochemical Approach to Identify Auxin Receptor

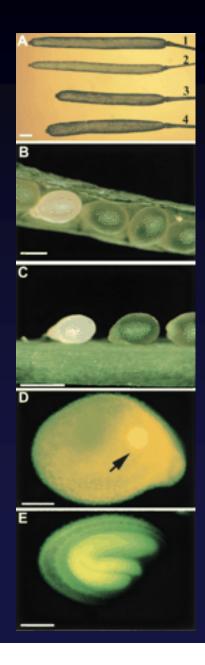
Isolation of auxin binding proteins

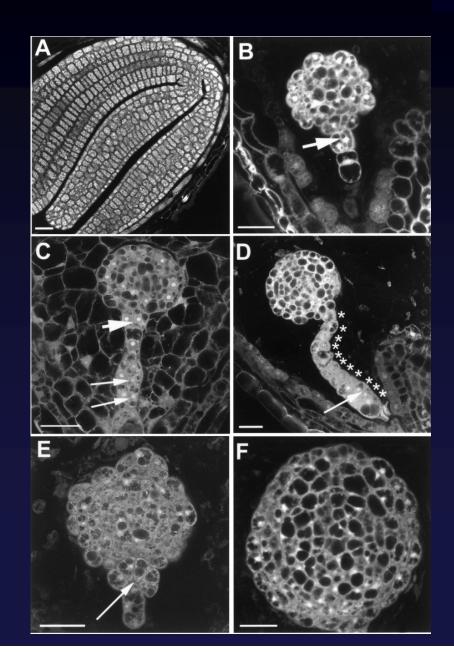
- Azidolabeling
- Affinity chromatography

Protein sequencing, cDNA screening, gene identification

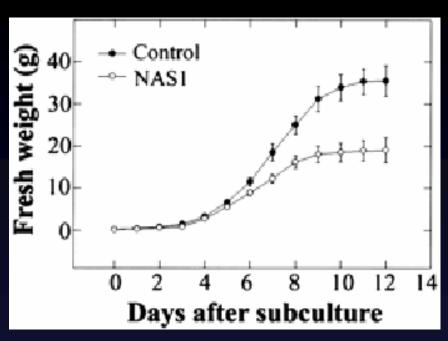
=> Auxin Binding Protein (ABP1)

Reverse Genetic – Embryo Lethal abp1 Mutant



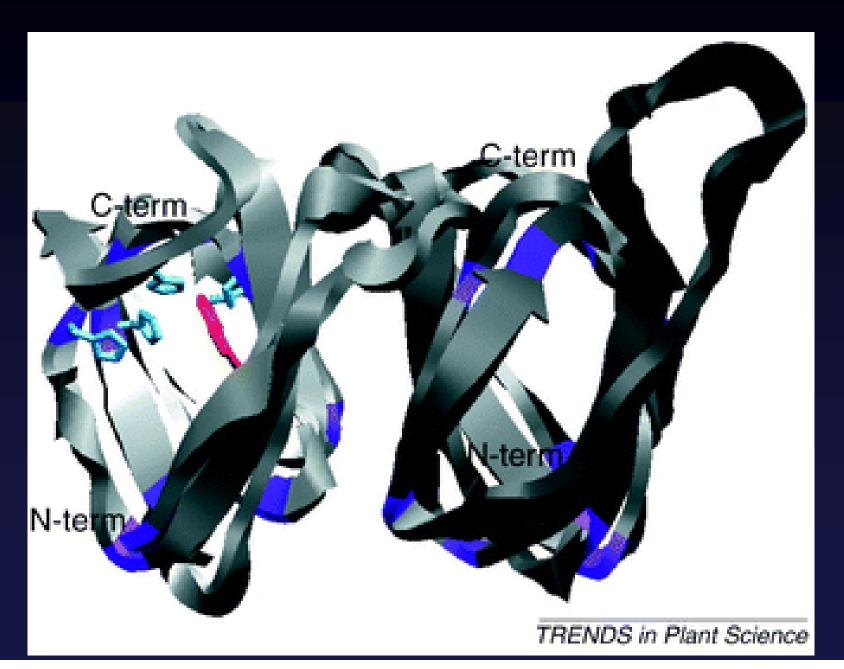


ABP1 Antisense
BY-2 Cells Display
Defects
in Auxin Dependent
Cell Elongation

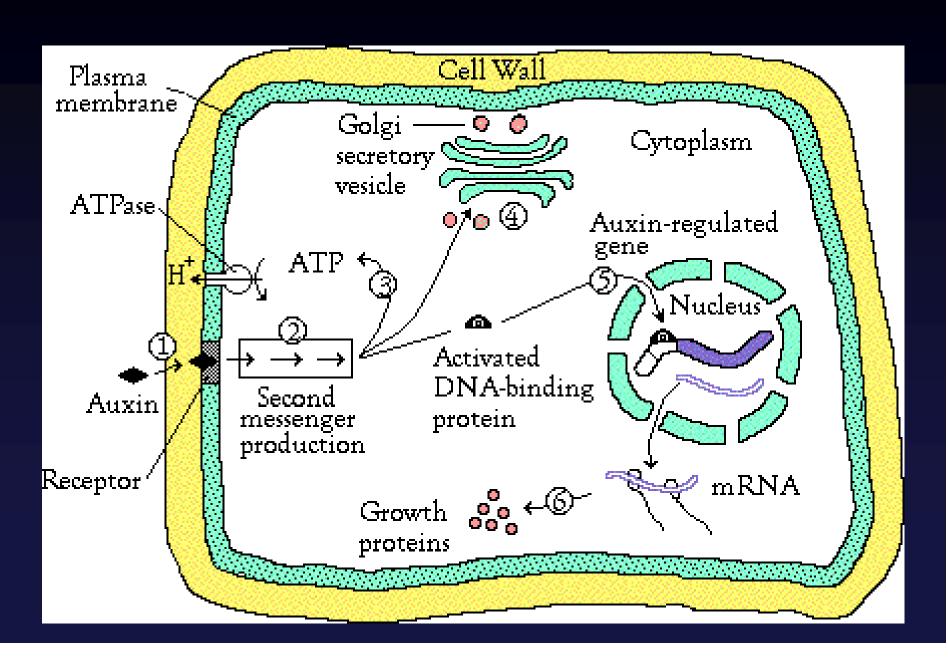




ABP1 – Structure



Optimistic Model for ABP1 Action



Genetic Approach to Identify Auxin Receptor

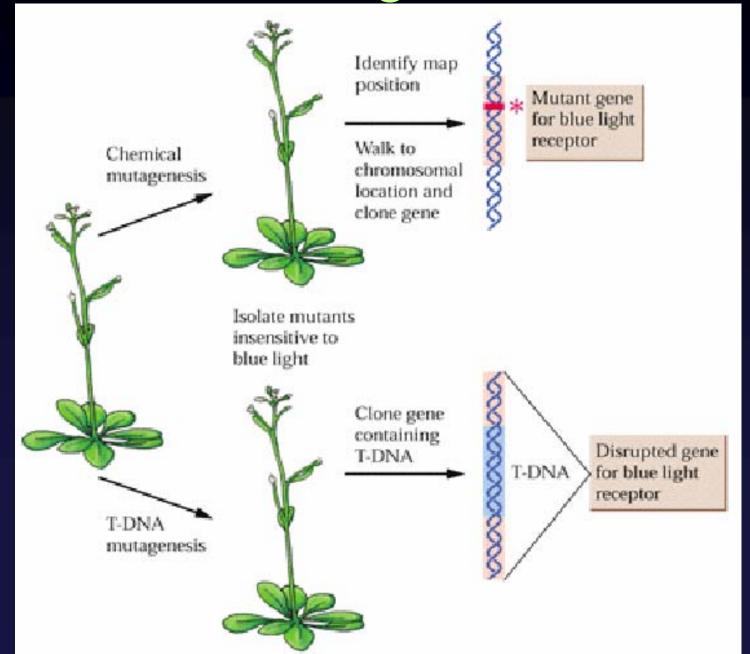
- Auxin resistant (axr): axr1 axr6
- Transport inhibitor response (tir): tir1 tir7

Morphological mutants (monopteros, bodenlos, etc.)

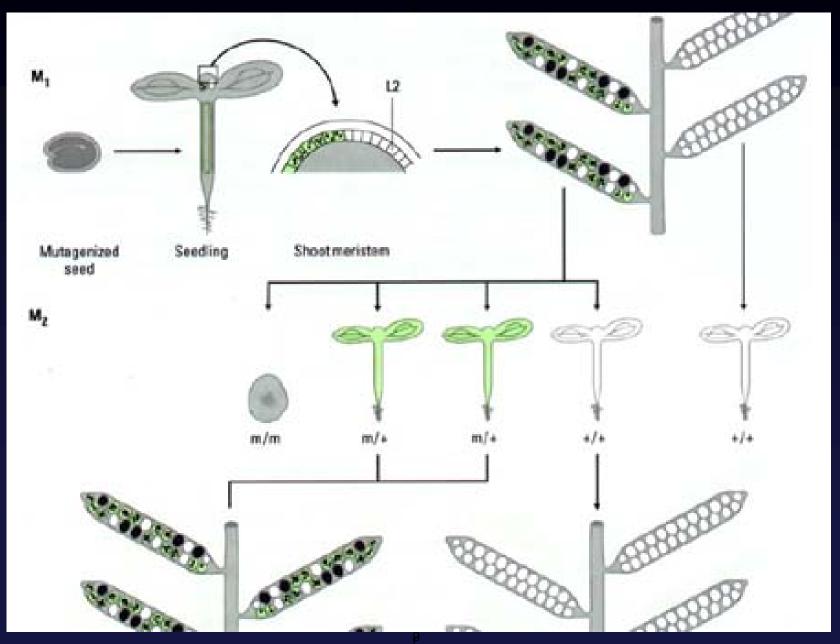
=> Role of regulated protein degradation and transcriptional regulation in auxin signaling

None of the identified gene looks like a receptor

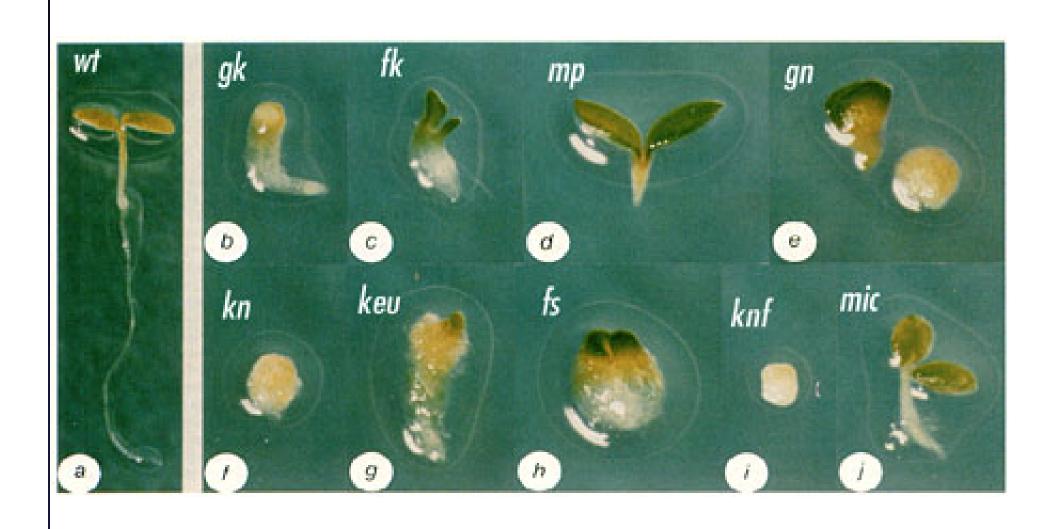
Forward genetics



EMS Mutagenesis



Mutant Screen at Seedling Level



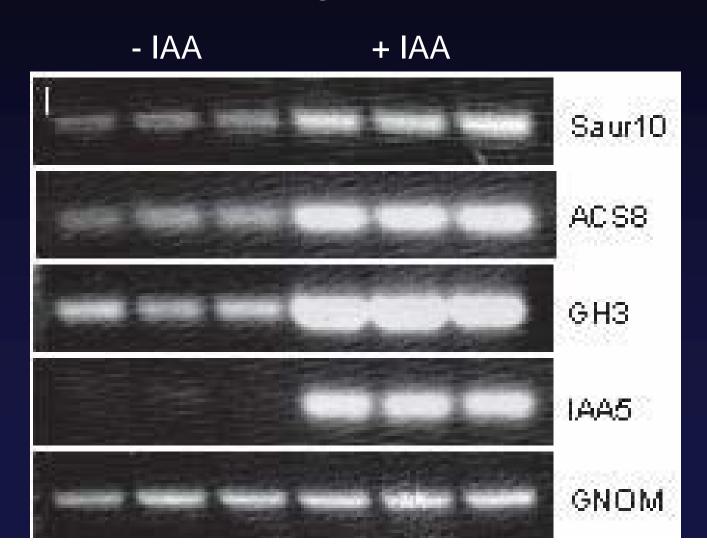
Molecular Biology Approach to Elucidate Auxin Signaling

Does auxin regulate gene expression?

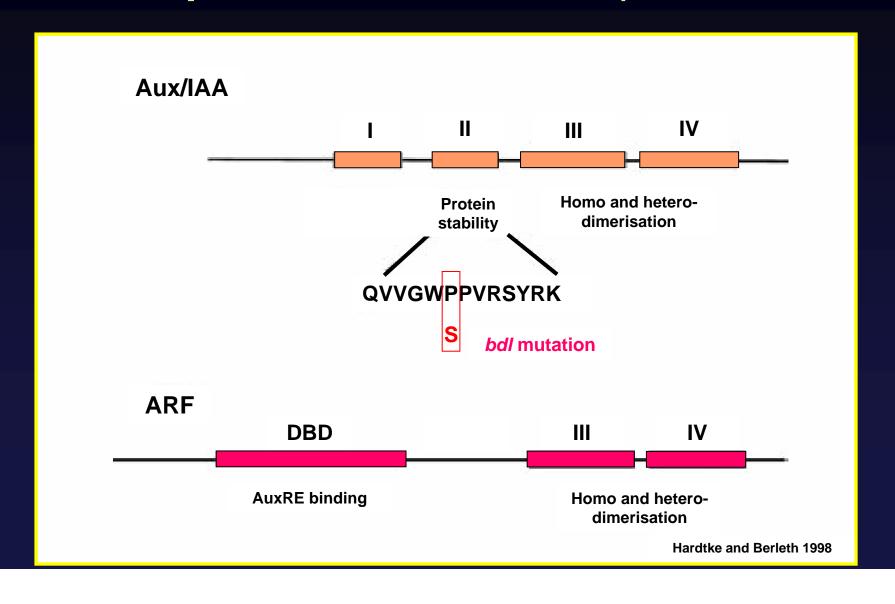
- Rapidly upregulated mRNAs (GH3, SAUR, AUX/IAA genes)
- One hybrid screen with Auxin Response Elements
 => Auxin Response Factors (ARF)
- Two hybrid => AUX/IAAs interact with ARFs

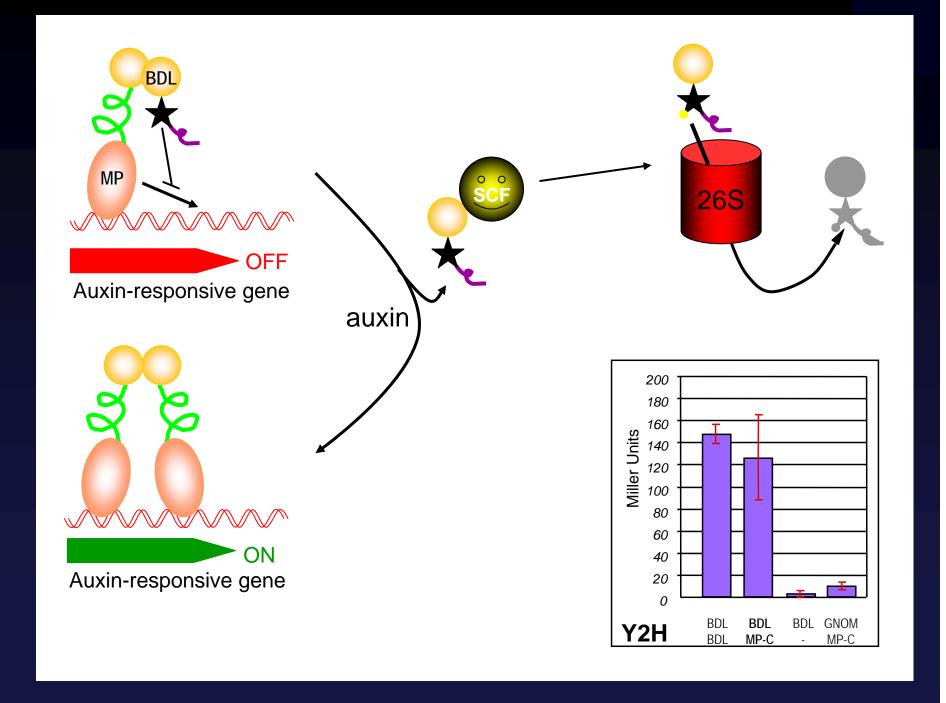
Molecular Biology Approach to Elucidate Auxin Signaling

RT-PCR

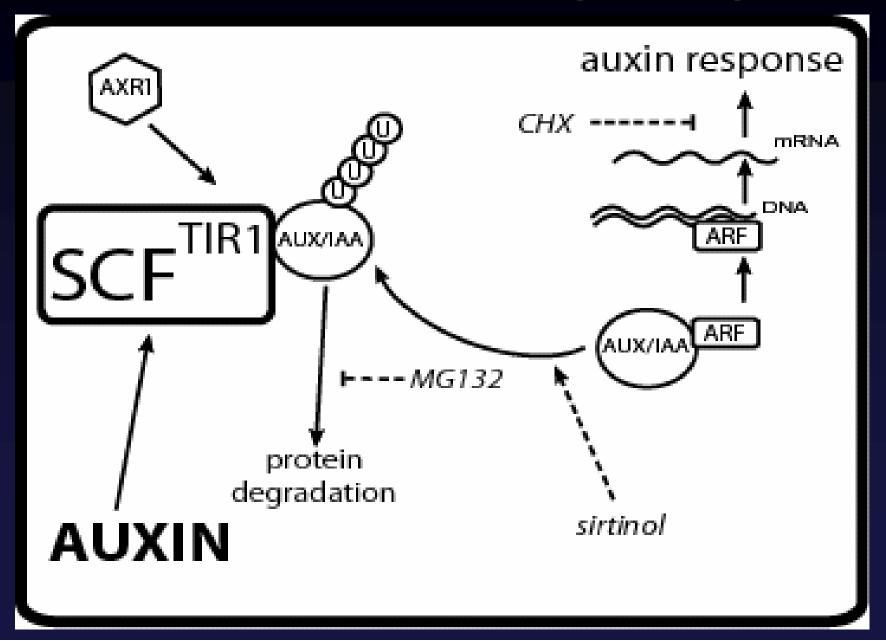


Some ARFs are **Activators**, whereas Aux/IAA **Repressors** of Auxin Response





Genomic Auxin Signaling



Summary for Auxin Signaling

Biochemical approach – auxin binding protein ABP1

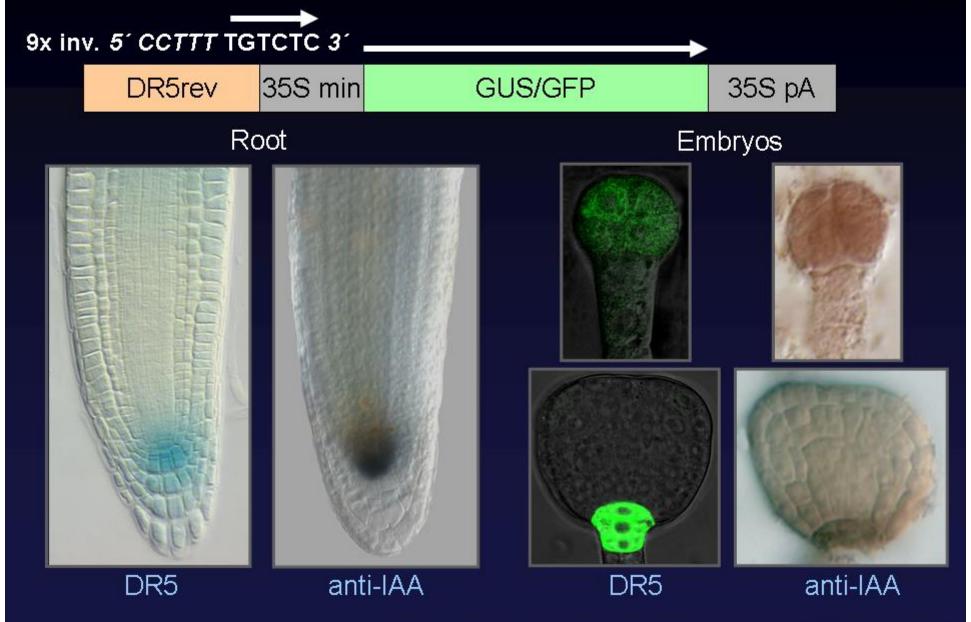
binds auxin, important in embryogenesis, precise role unclear

Genetic approach – role of protein degradation (axr1, tir1)

Molecular approach – auxin regulates expression ARE in promotors of auxin regulated genes ARF transcription factors binds to ARE AUX/IAA proteins repress ARF and are degraded upon auxin signal

DR5 Auxin Response Reporter

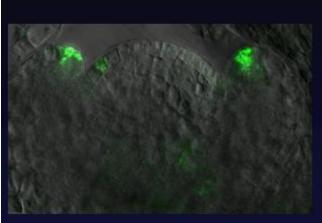


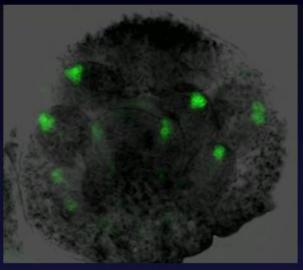


Ulmasov et al. 97; Sabatini et al. 99; Benková et al. 03; Friml et al. 03

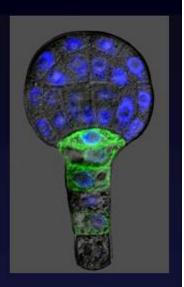
Local Auxin Gradients Require Active Polar Auxin Transport

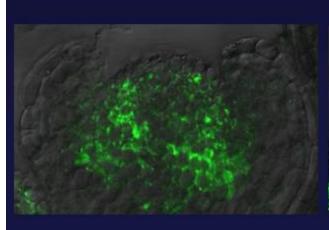


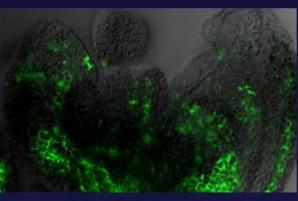




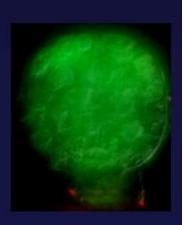












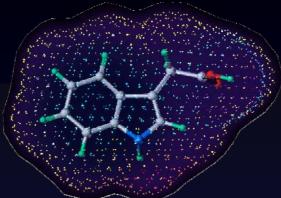
Auxin Transport

Proteins involved in auxin transport

- -PIN proteins (efflux)
- -AUX1 proteins (influx)

AUXIN TRANSPORT

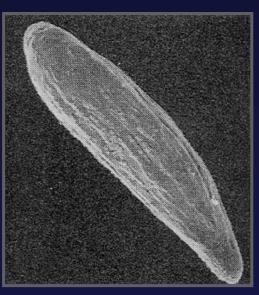
mediates



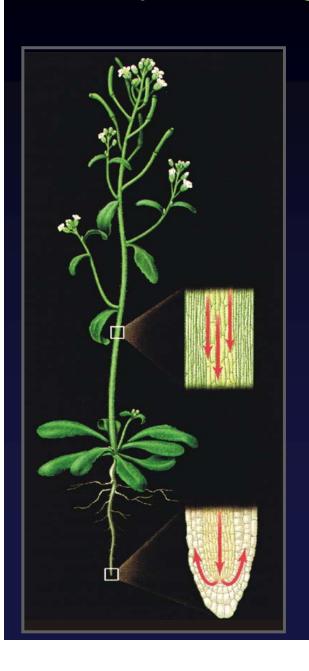
embryos

- Embryo development
- Organ initiation and positioning
- Vascular tissue differentiation
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- Growth responses to light and gravity
- Apical hook formation

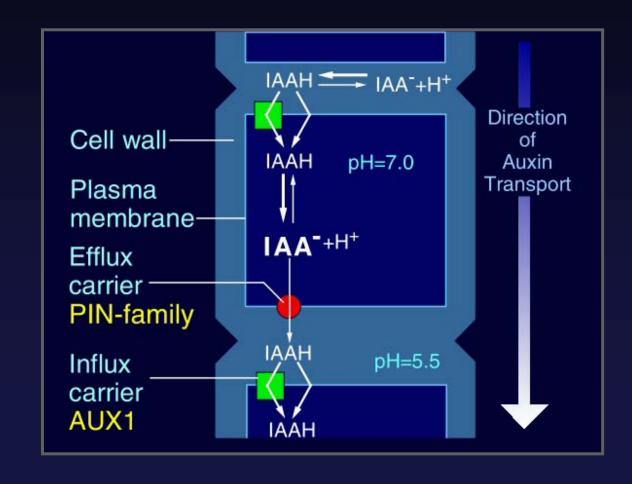




Physiology of Auxin Transport



Chemiosmotic hypothesis

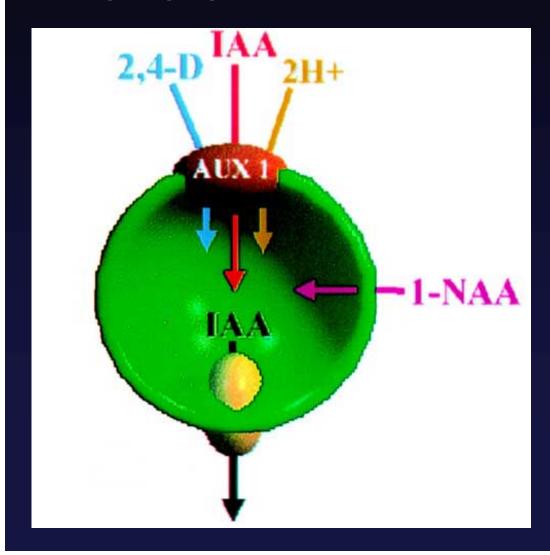


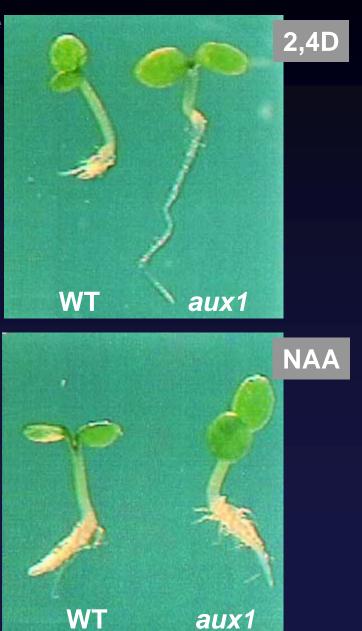
Auxin Influx

aux1 is Resistant to Auxin

aux1 phenotype

Transport properties of different auxins





NAA Rescues aux1 Phenotype

- NAA



+ NAA



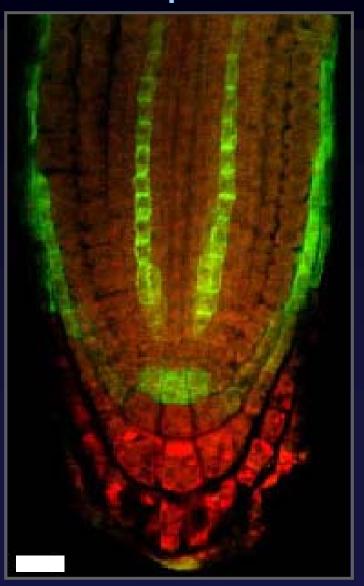
AUX1 – Expression and Localization

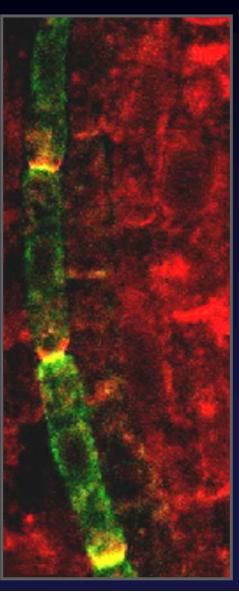
AUX1::GUS

AUX1 protein

PIN1/AUX1





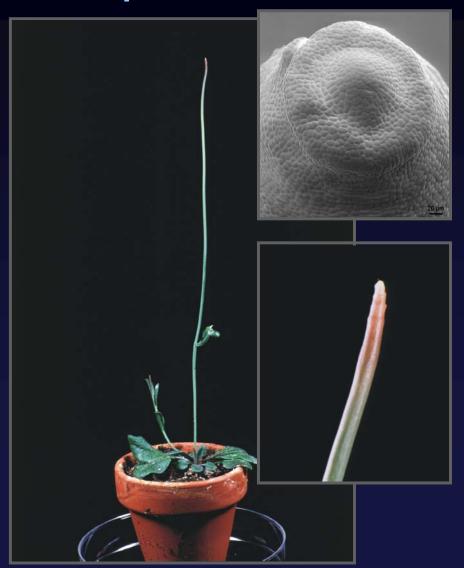


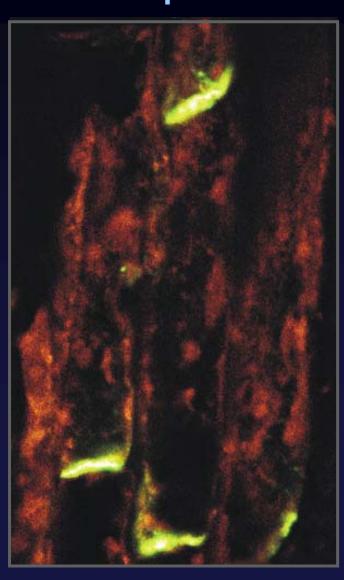
Auxin Efflux

PIN1 – the Auxin Efflux Carrier?

pin1 mutant

PIN1 protein



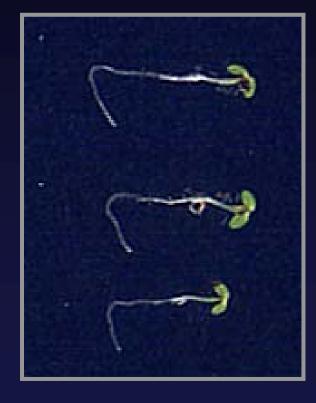


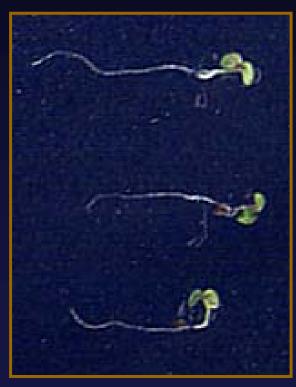
PIN2 – Root Gravitropism

PIN2 protein





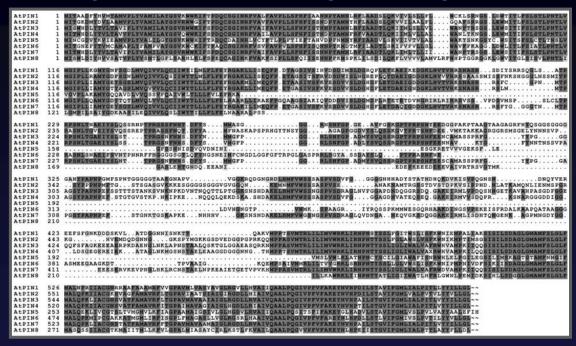




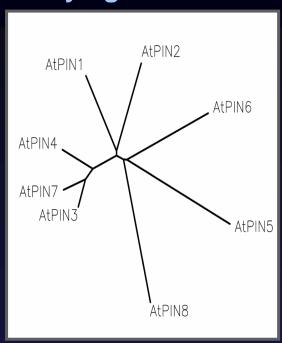


The Arabidopsis PIN Gene Family

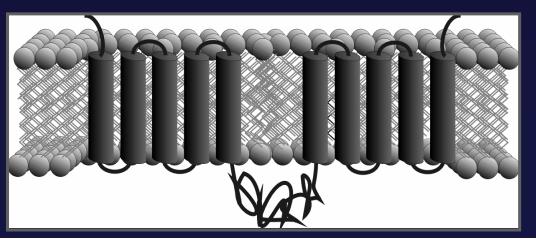
Comparison of *Arabidopsis* PIN proteins



Phylogenetic tree



Membrane topology model



What is Molecular Role

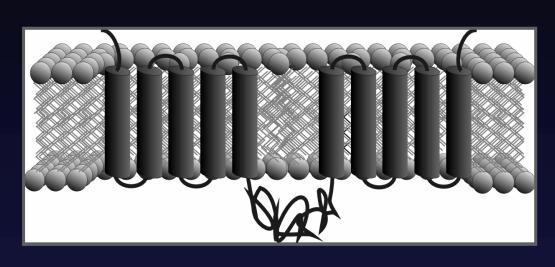
of PIN Proteins

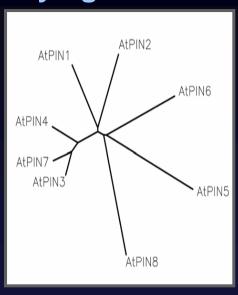
in Auxin Transport?

PINs Are Essential Components of Auxin Transport

Putative topology of PIN proteins

Phylogenetic tree



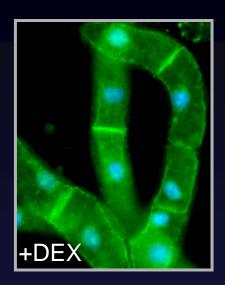


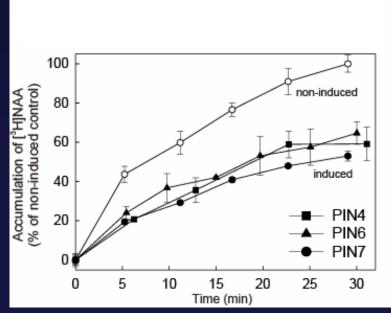
- All defects in *pin* loss-off-function mutants are in auxin transport-dependent processes and can be phenocopied by auxin transport inhibitors
- Local auxin distribution (gradients) are affected in pins
- Polar PIN localization determines direction of auxin flow

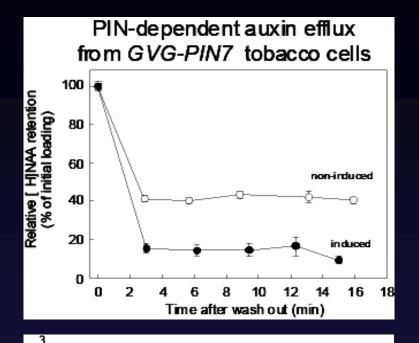
PINs Are Rate-limiting Factors in Auxin Efflux

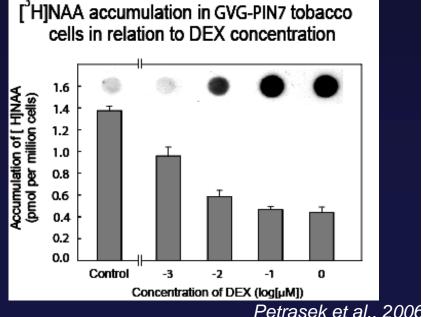
Inducible PIN1 expression



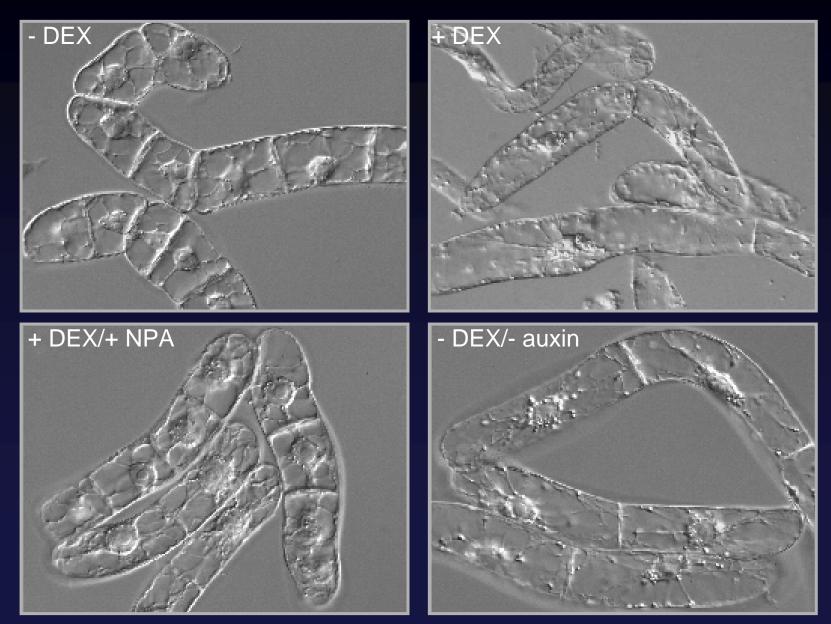






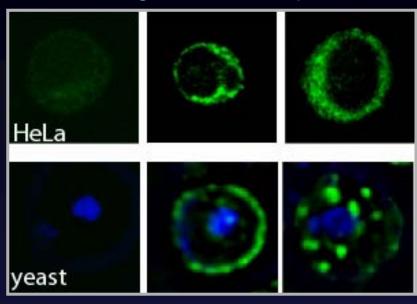


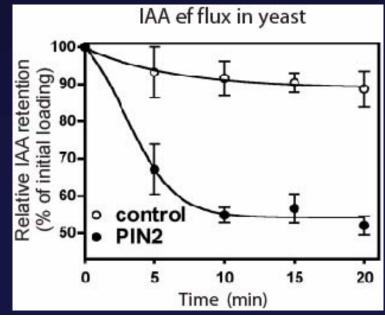
PIN-induced Phenotypes in BY-2 Cells



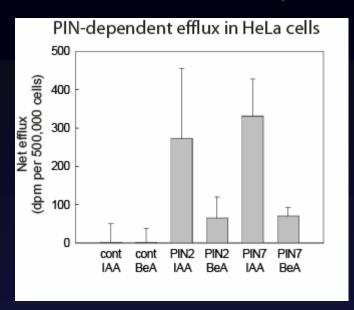
Expression of PINs in HeLa and Yeast

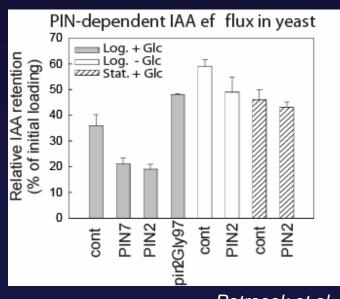
Heterologous PIN2 expression





auxin efflux activity





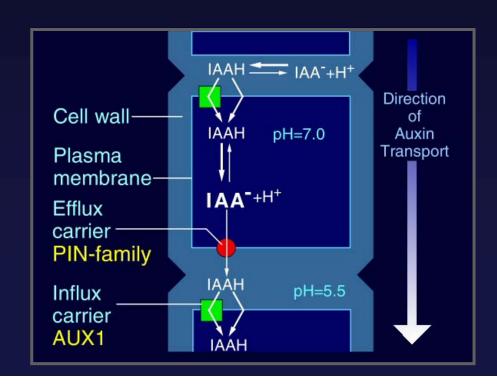
Petrasek et al., 2006

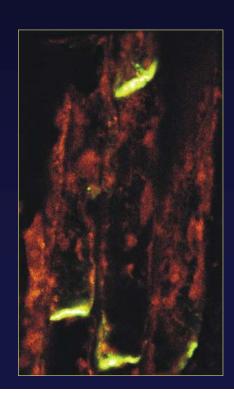
PIN proteins are rate-limiting factors in auxin efflux from cells

and

the polarity of their subcellular localization determines direction of intercellular auxin flow

Cellular Polarity of PIN Localization and Directionality of Intercellular Auxin Flow



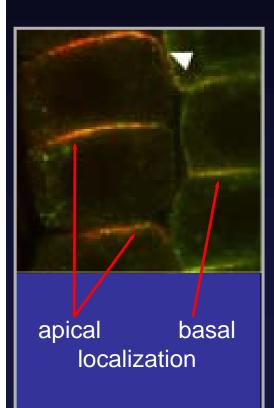


PIN-specific Signals for Polar Targeting

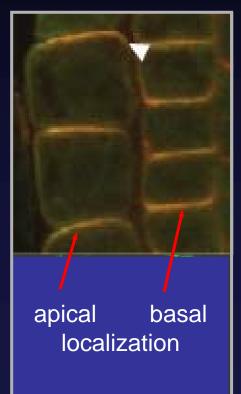
PIN2pr::PIN2:HA

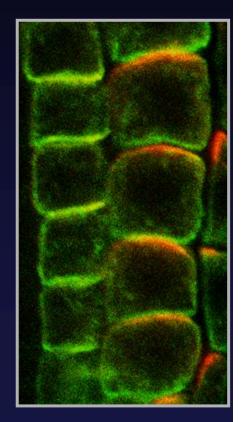
PIN2pr::PIN1:HA PIN2pr::PIN1:GFP

PIN1/PIN2





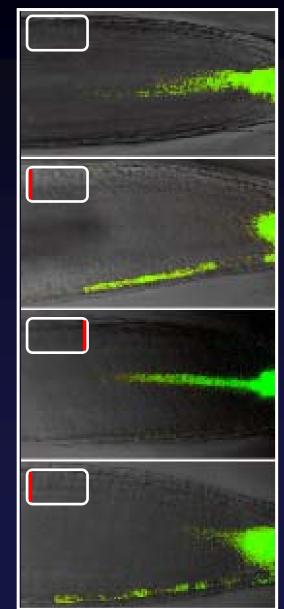




PIN Polarity Determines Direction

DR5rev::GFP

of Auxin Flow gravitropism

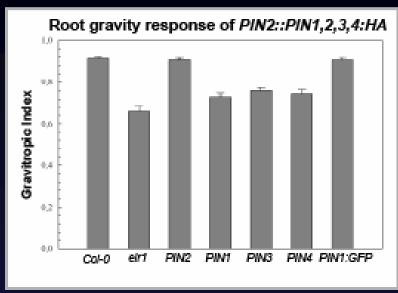


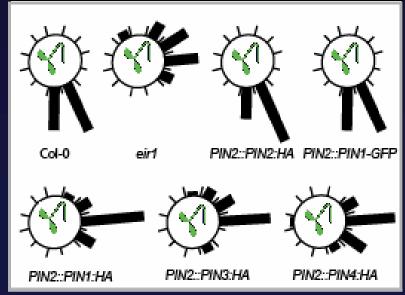
eir1

PIN2::PIN2:HA

PIN2::PIN1:HA
PIN2::PIN1:GFP-2

PIN2::PIN1:GFP-3





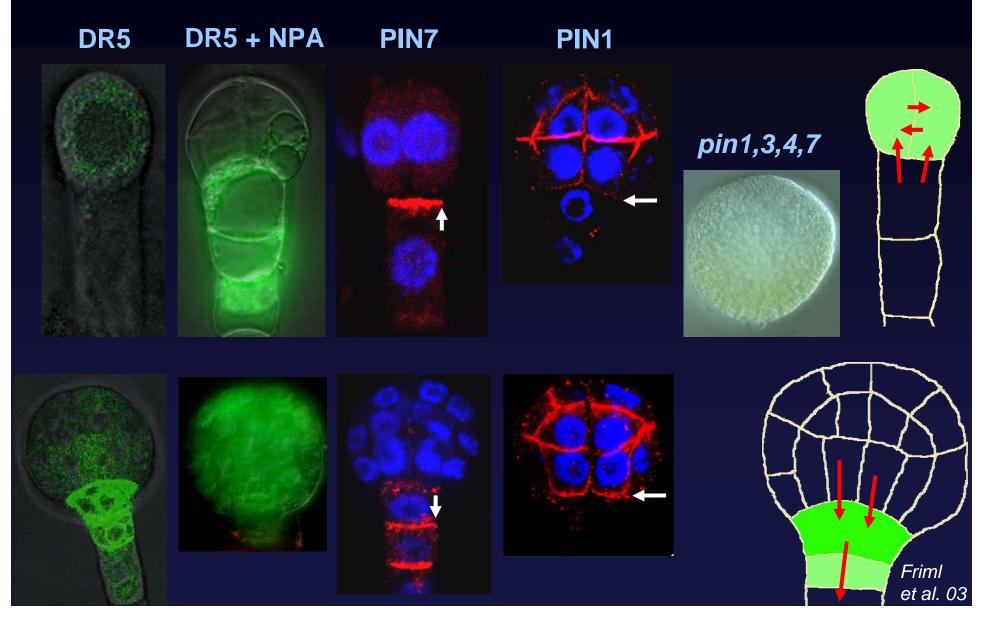
Wisniewska et al., 2006

PIN proteins are rate-limiting factors in auxin efflux from cells

and

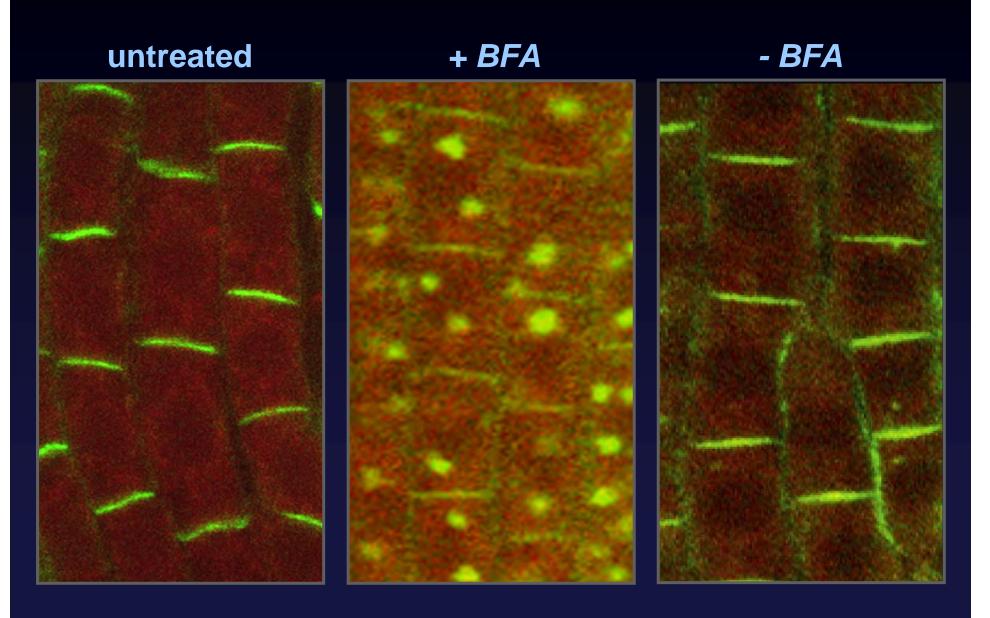
the polarity of their subcellular localization determines direction of intercellular auxin flow

Auxin in Embryonic Apical-Basal Axis Formation

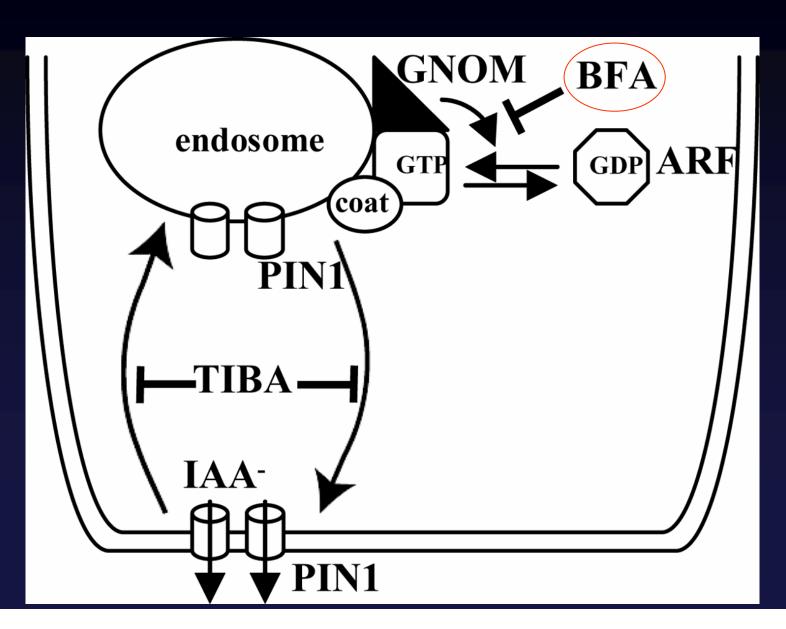


Constitutive Cycling of PINs

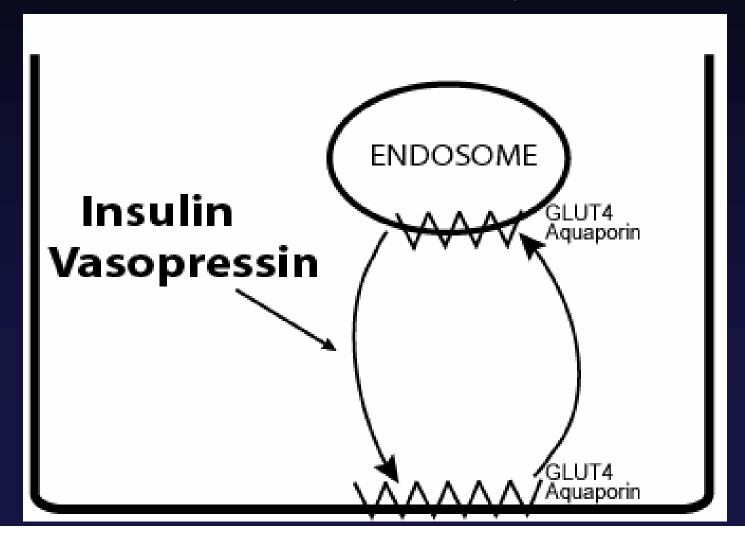
PIN1 Subcellular Movement



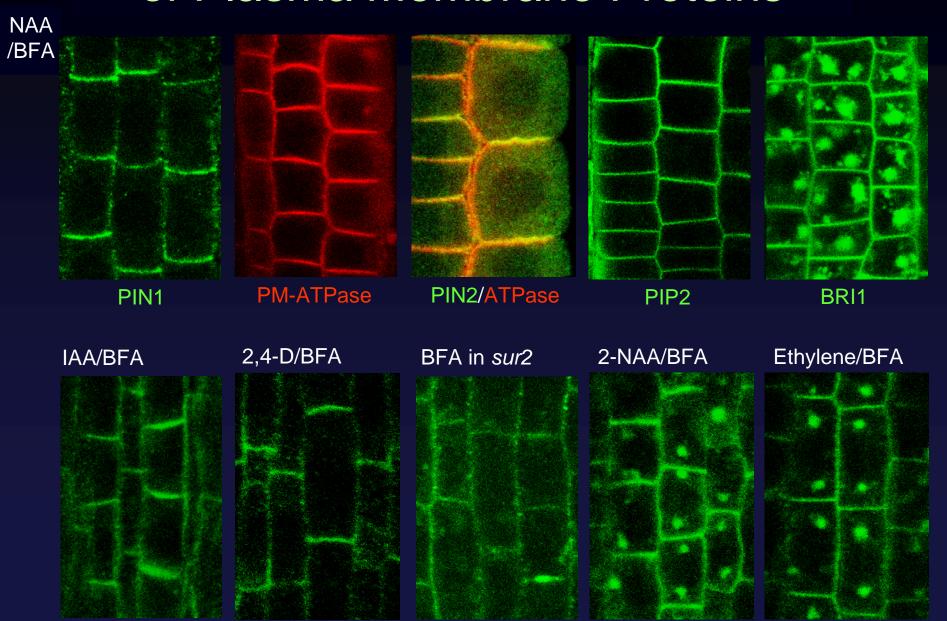
Dynamic Movement of PIN Proteins



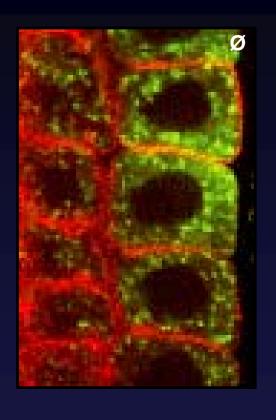
Subcellular Cycling – Means to Modulate Protein Activity?

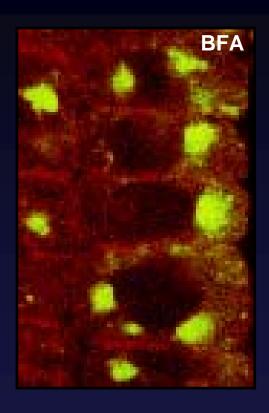


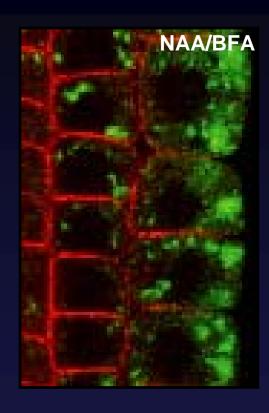
Auxin Inhibits Internalization of Plasma Membrane Proteins



Auxin Effect on Endosome Trafficking





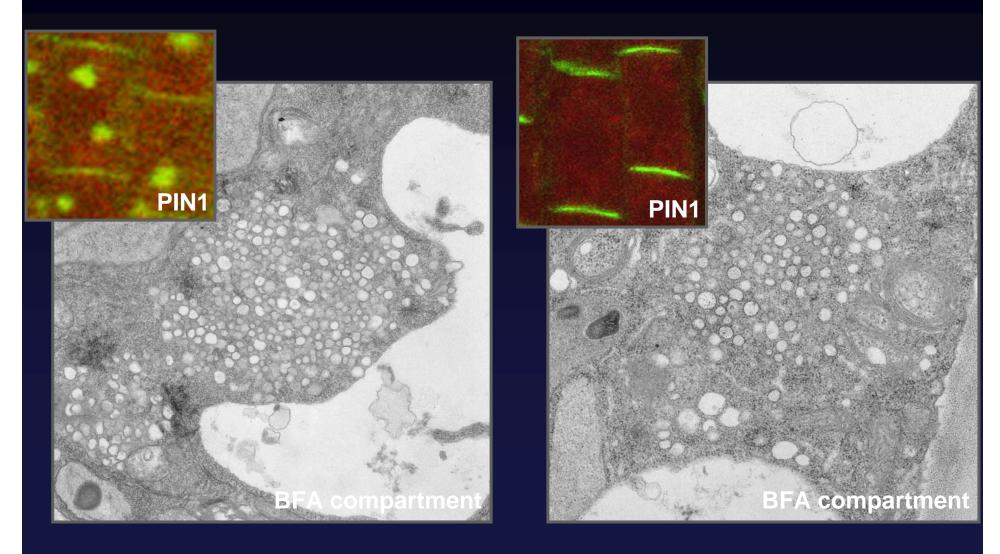


- endosoms
- H+ATPase

Place of Auxin Action in Protein Cycling

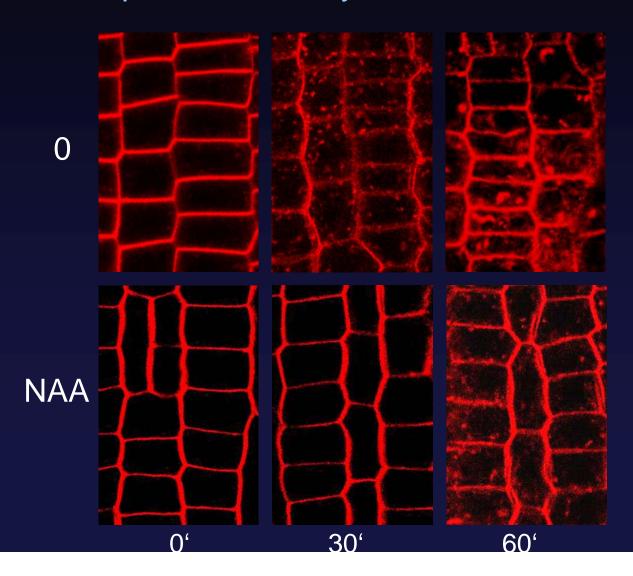
BFA

Auxin + BFA



Auxin Inhibits Endocytosis

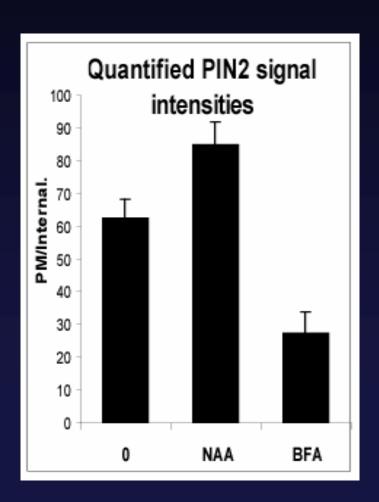
Uptake of endocytic tracer FM4-64



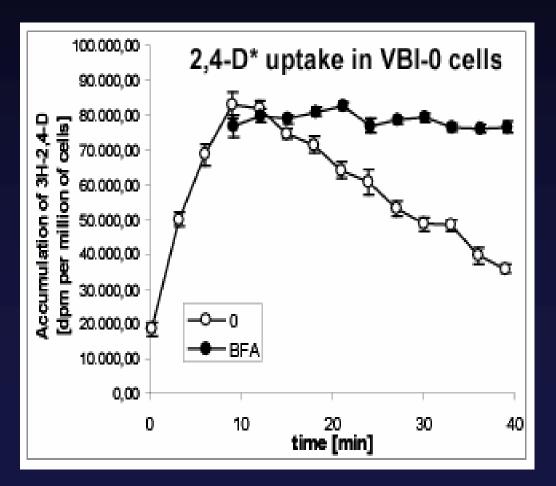
Paciorek et al.. 2005

Auxin Increases PIN Levels at Cell Surface and Stimulates its own Efflux

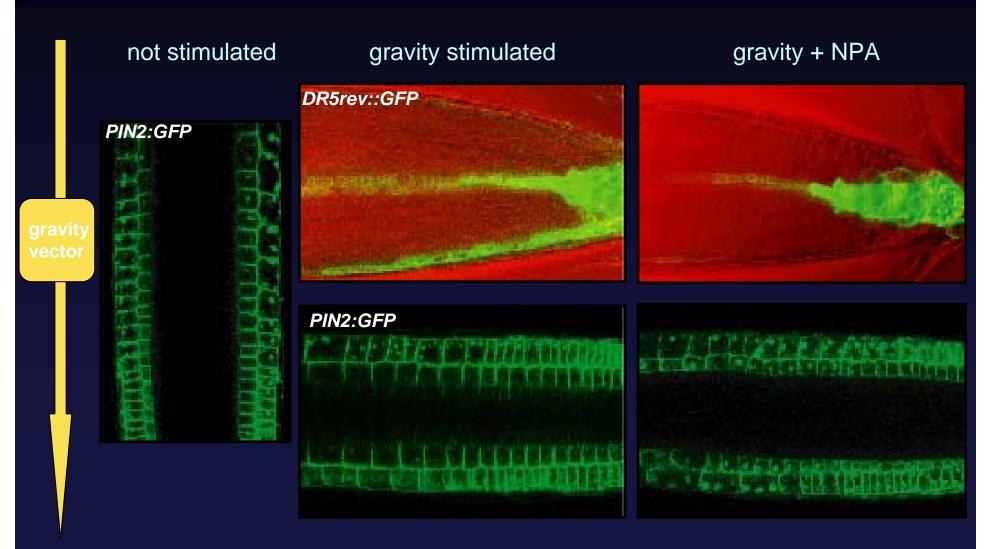
PIN2 levels at PM



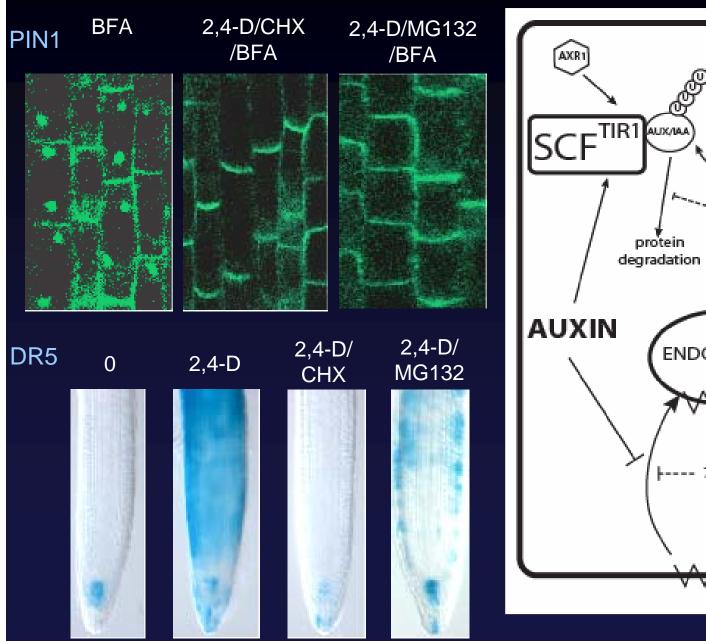
Auxin efflux in tobacco cells

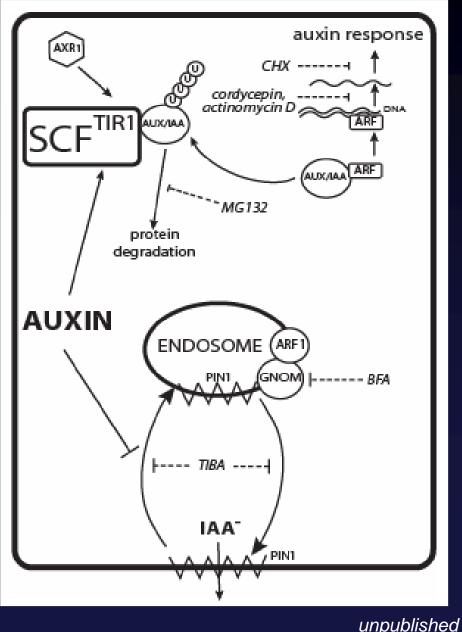


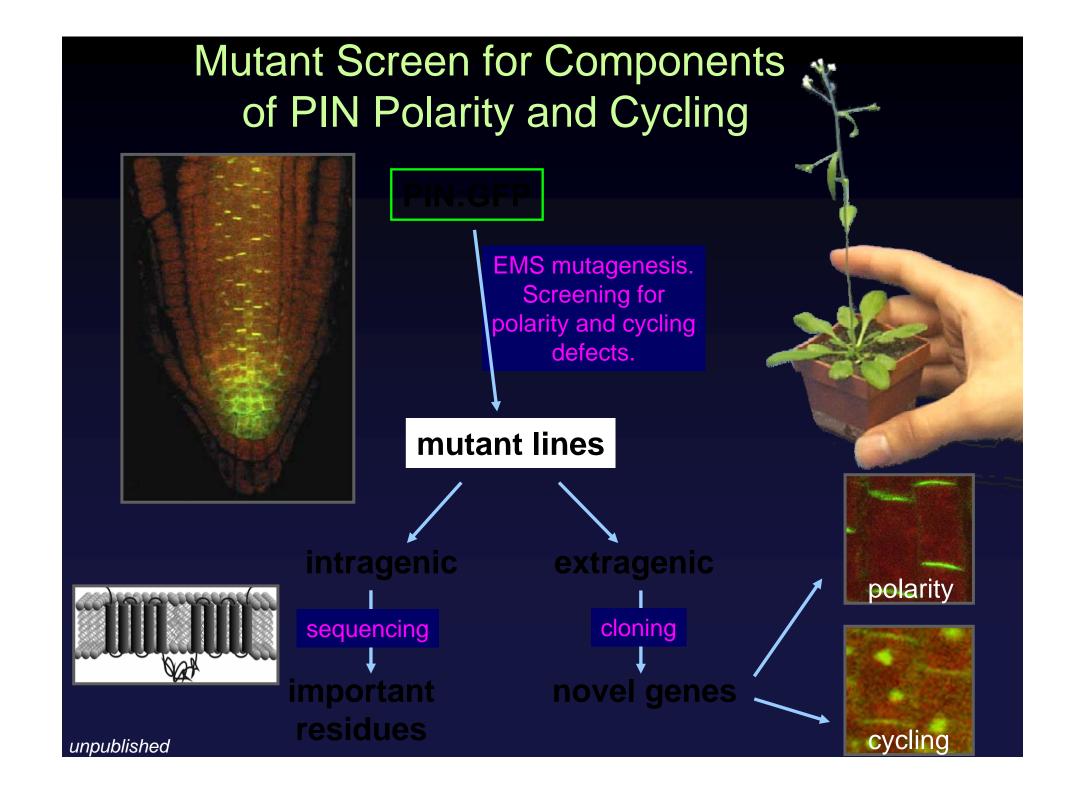
in planta Correlation between Cycling and Auxin Flow



Novel Pathway of Auxin Action



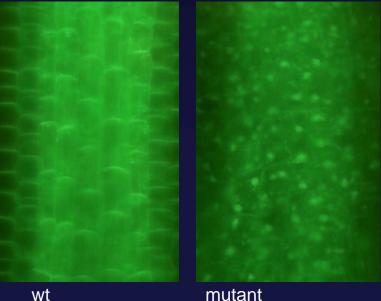




"Cell Biological" Mutant Screens in Progress:

Auxin effect on endocytosis: 3 confirmed mutants

30′ NAA 30 μM/90′BFA 50 μM



Auxin-resistant BFA patches mutants



Novel Pathway for Auxin Signaling

Auxin inhibits endocytosis including internalization of PIN proteins

This is mechanism by which auxin stabilizes PINs at the cell surface thus stimulating auxin efflux.

This auxin effect involves novel, genetically tractable auxin pathway