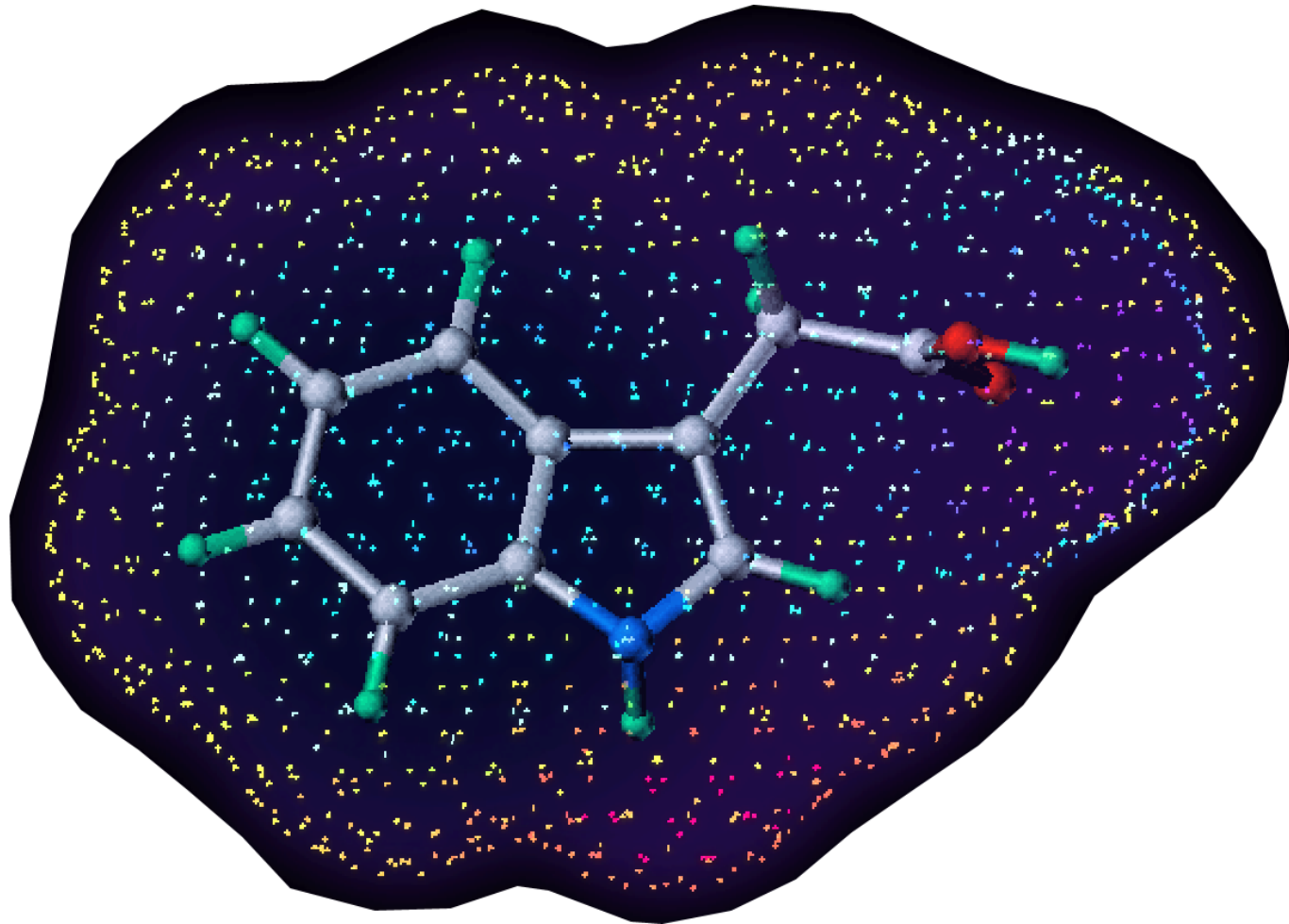
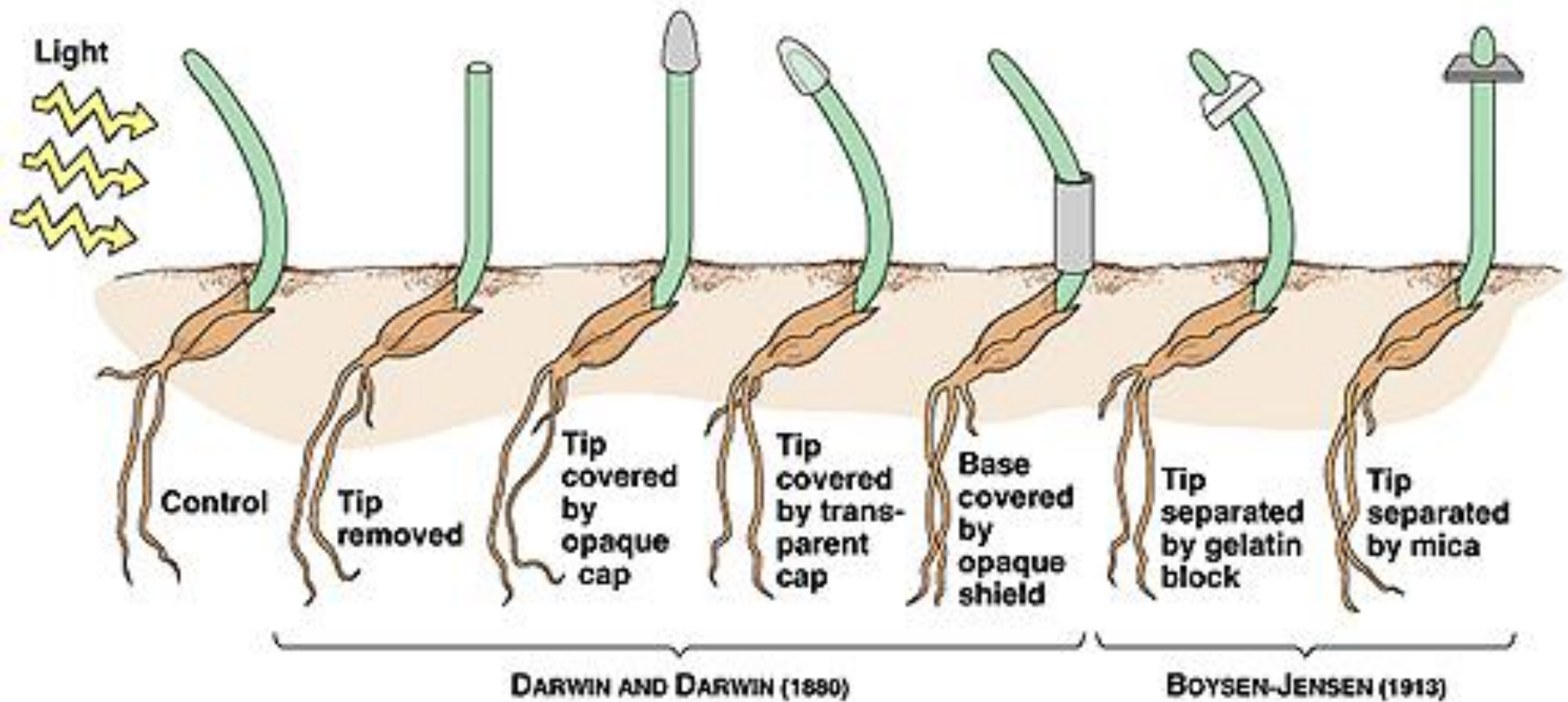


Auxin Signaling and Transport

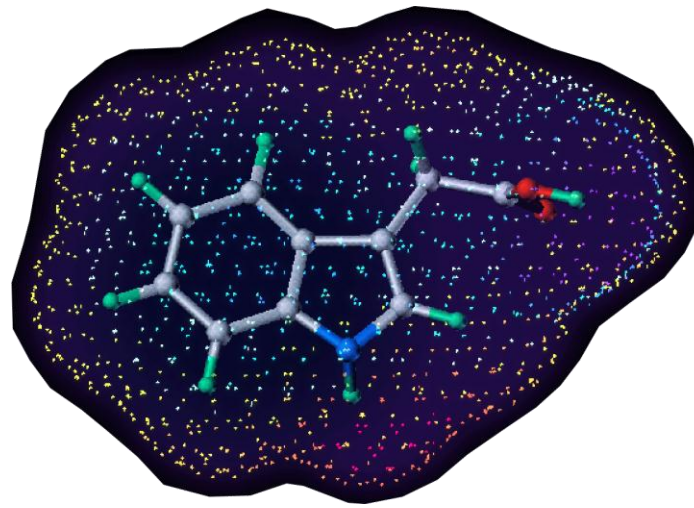


Discovery of the First Plant Signaling Molecule – Auxin and its Transport

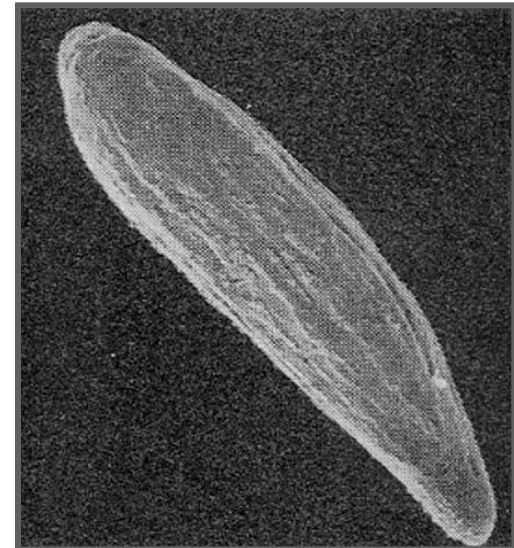
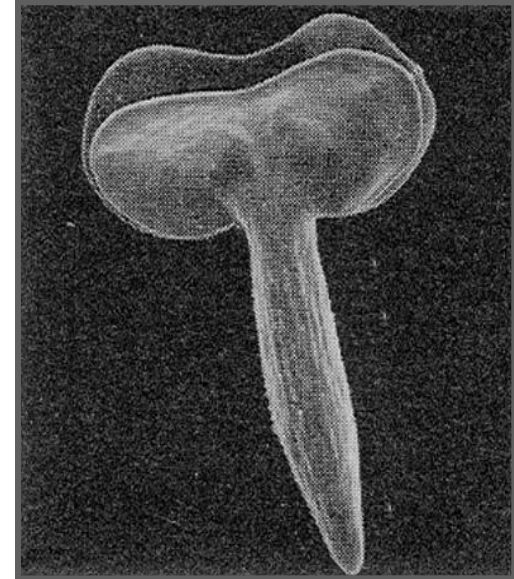


AUXIN

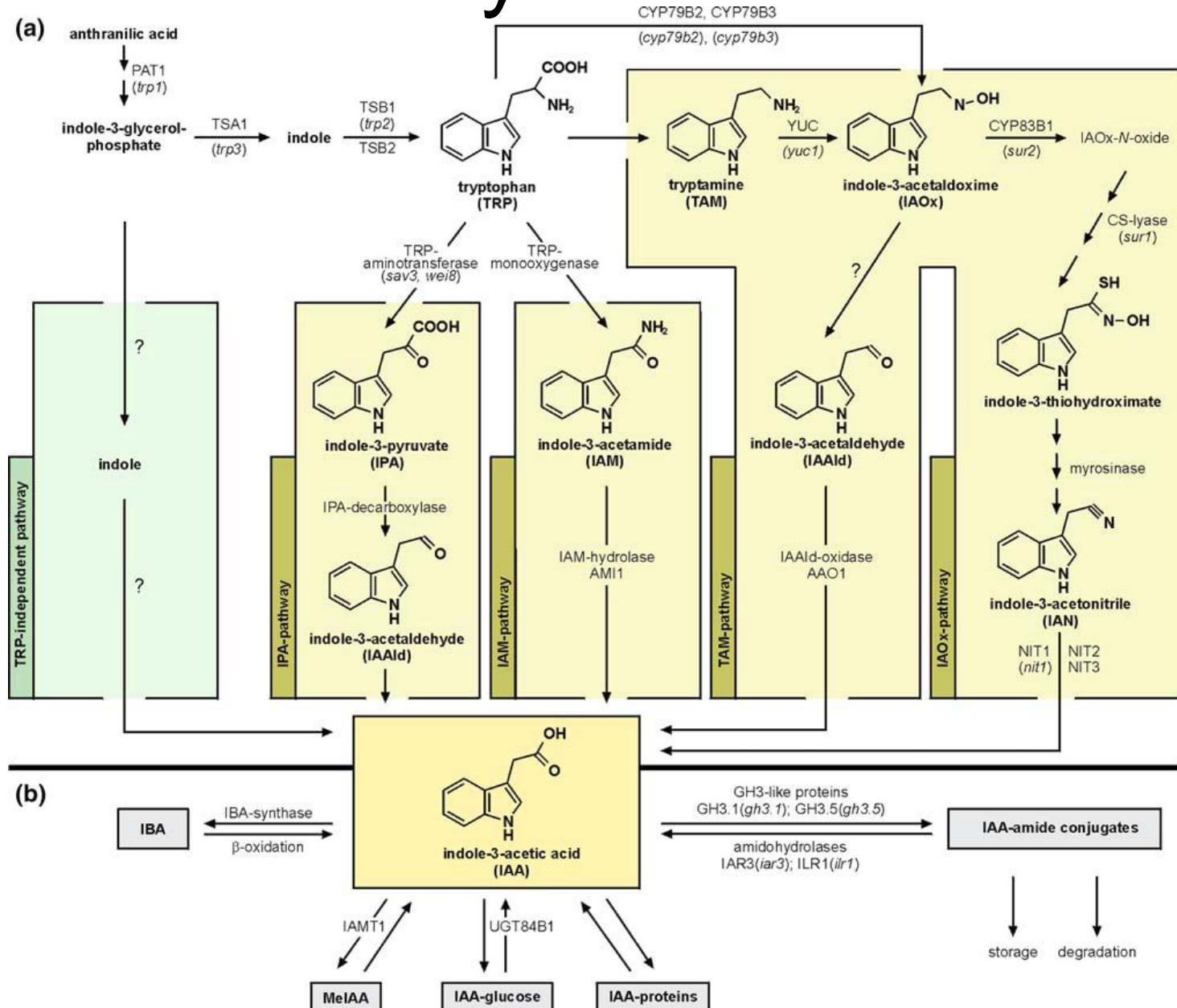
mediates



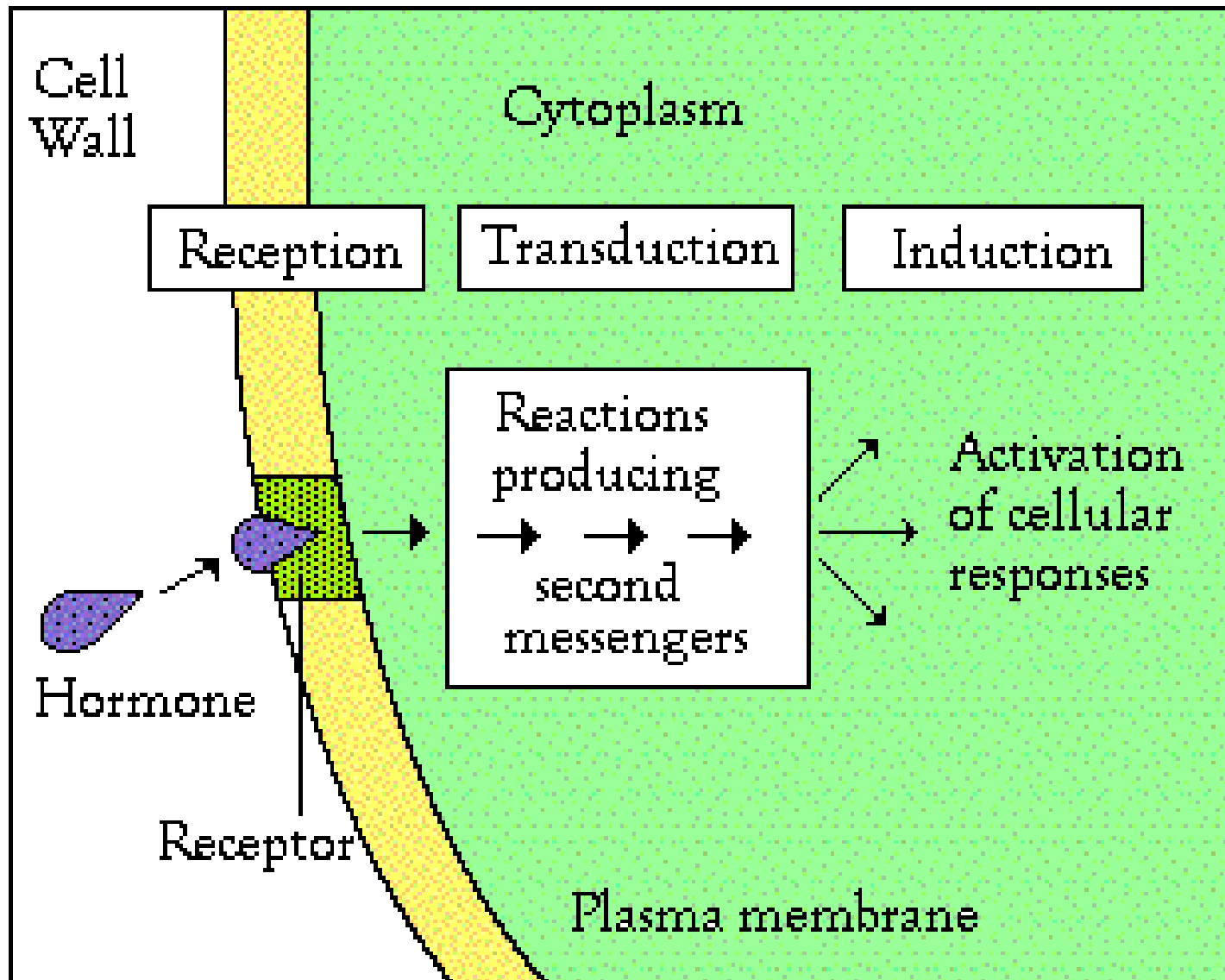
- Embryo development
- Organ initiation and positioning
- Vascular tissue differentiation
- Shoot and root elongation
- Growth responses to light and gravity
- Apical hook formation



Auxin biosynthesis



Signal Transduction



Biochemical Approach to Identify Auxin Receptor

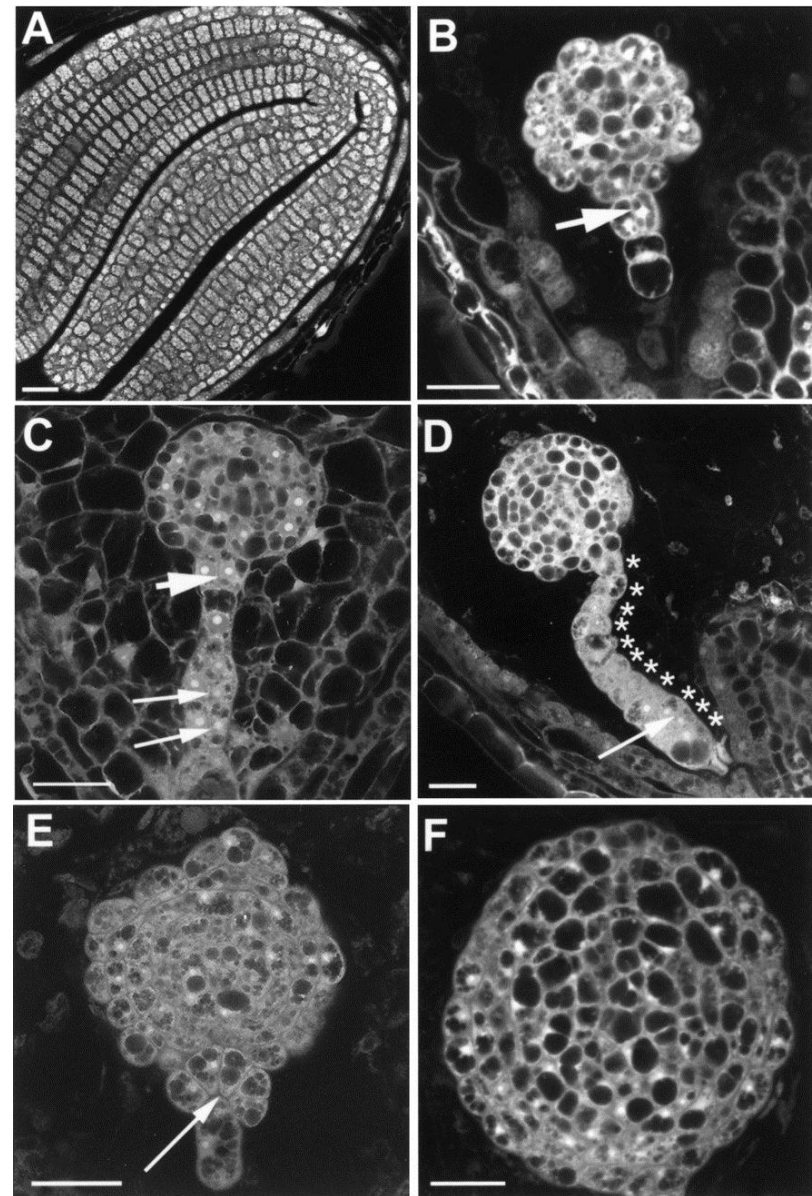
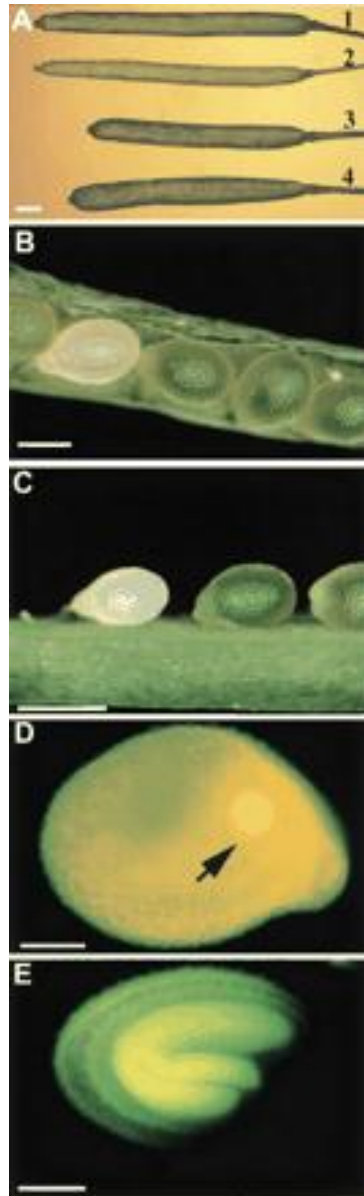
Isolation of auxin binding proteins

- Azidolabeling
- Affinity chromatography

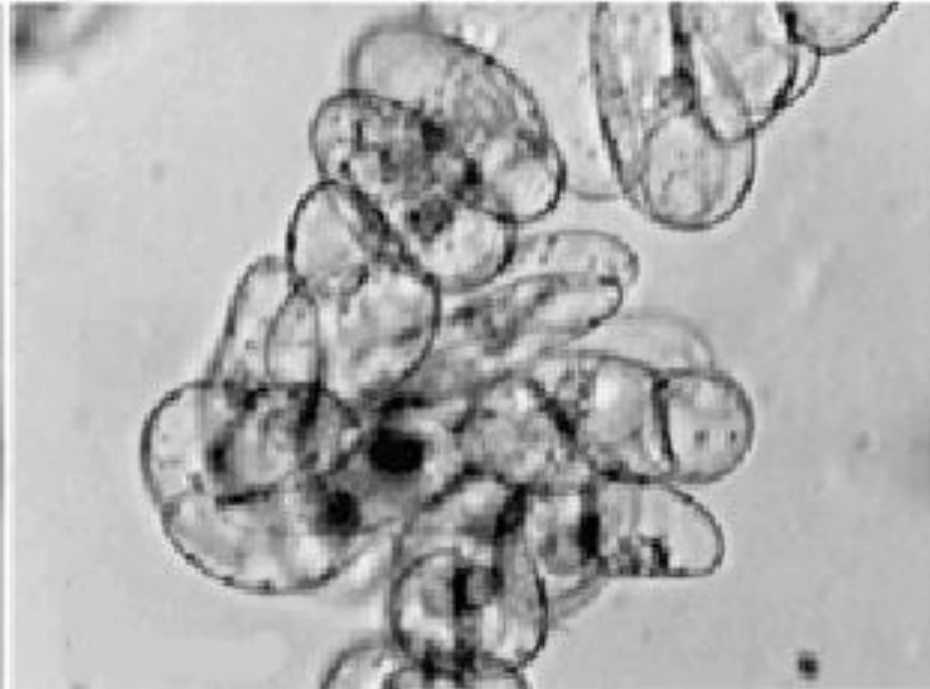
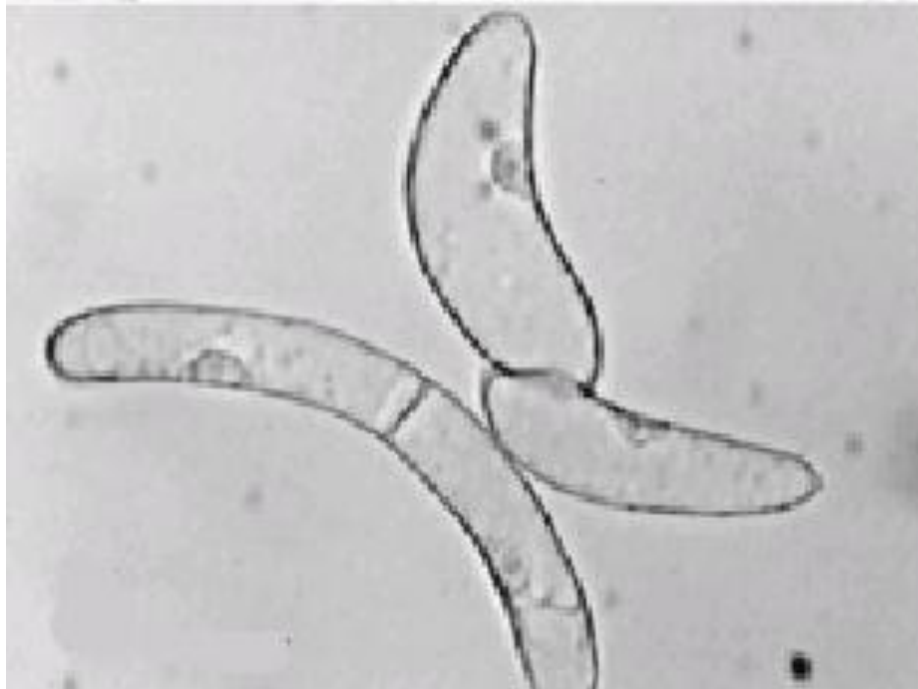
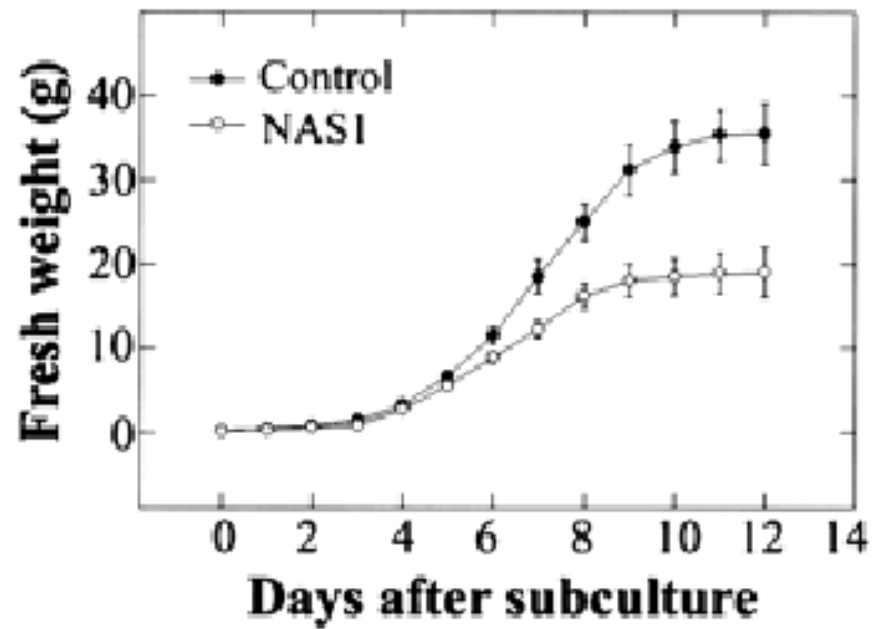
Protein sequencing, cDNA screening, gene identification

=> Auxin Binding Protein (ABP1)

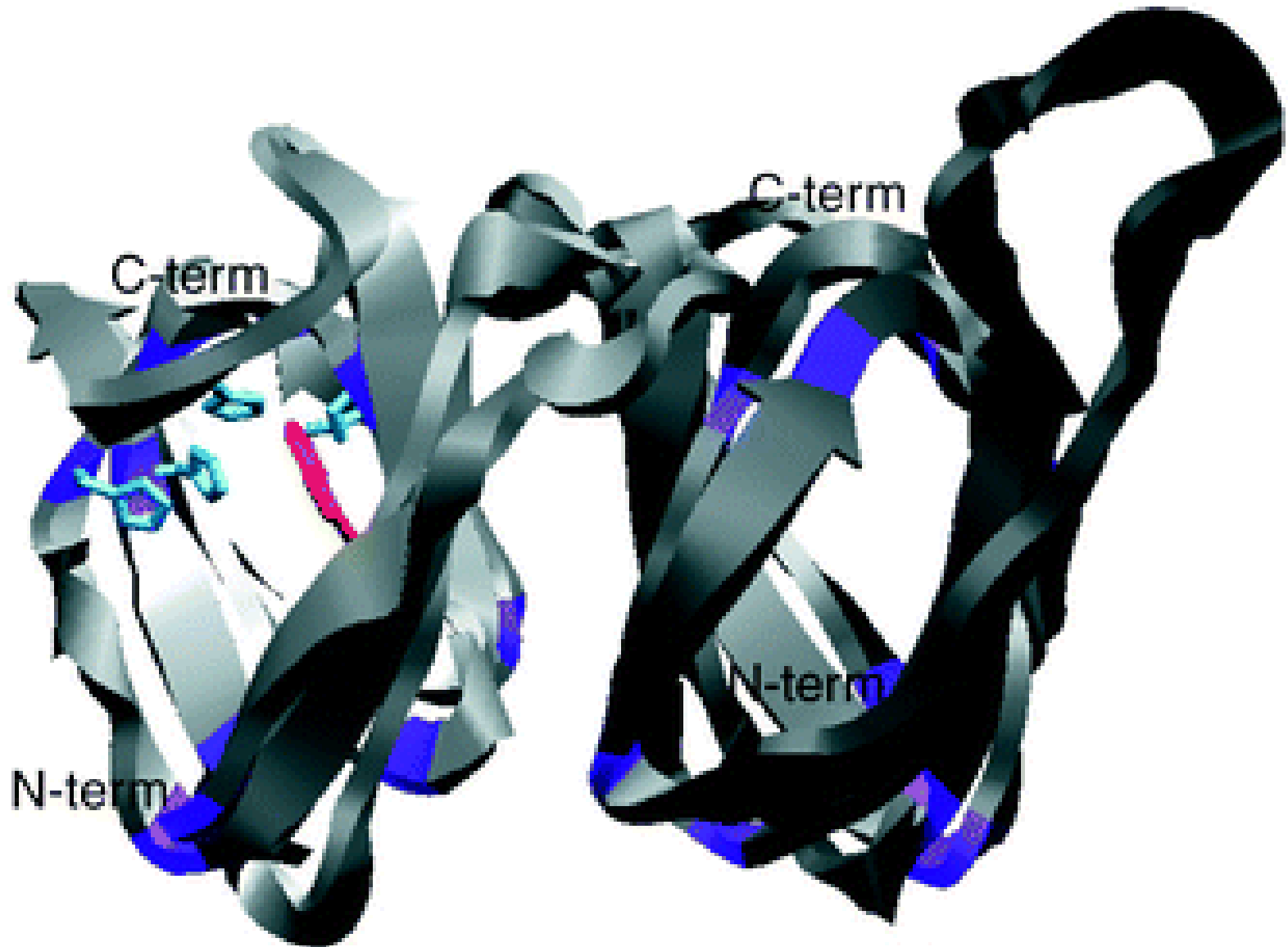
Reverse Genetic – Embryo Lethal *abp1* Mutant



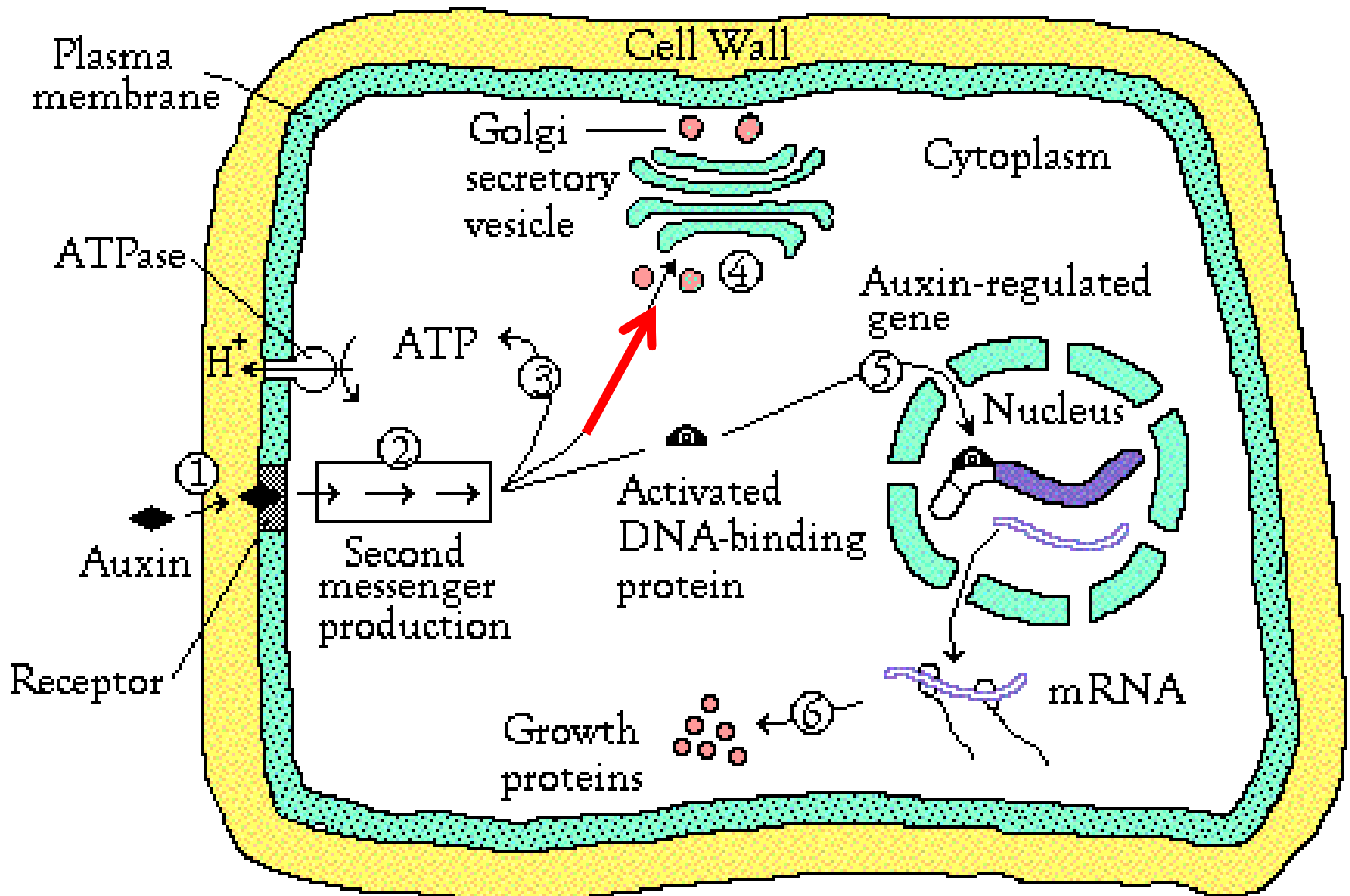
ABP1 Antisense BY-2 Cells Display Defects in Auxin Dependent Cell Elongation



ABP1 – Structure



Optimistic Model for ABP1 Action



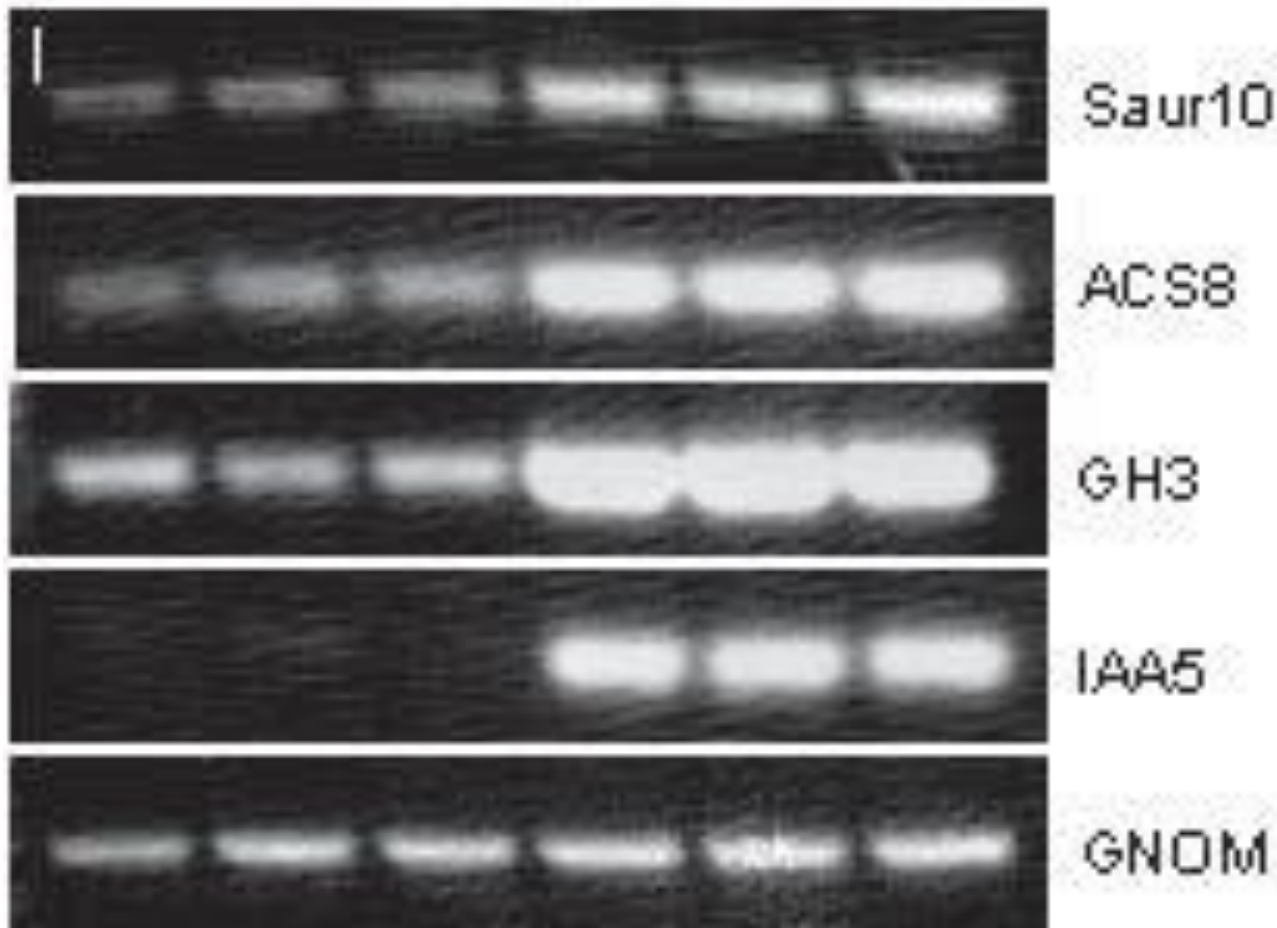
Molecular Biology Approach to Elucidate Auxin Signaling

Does auxin regulate gene expression?

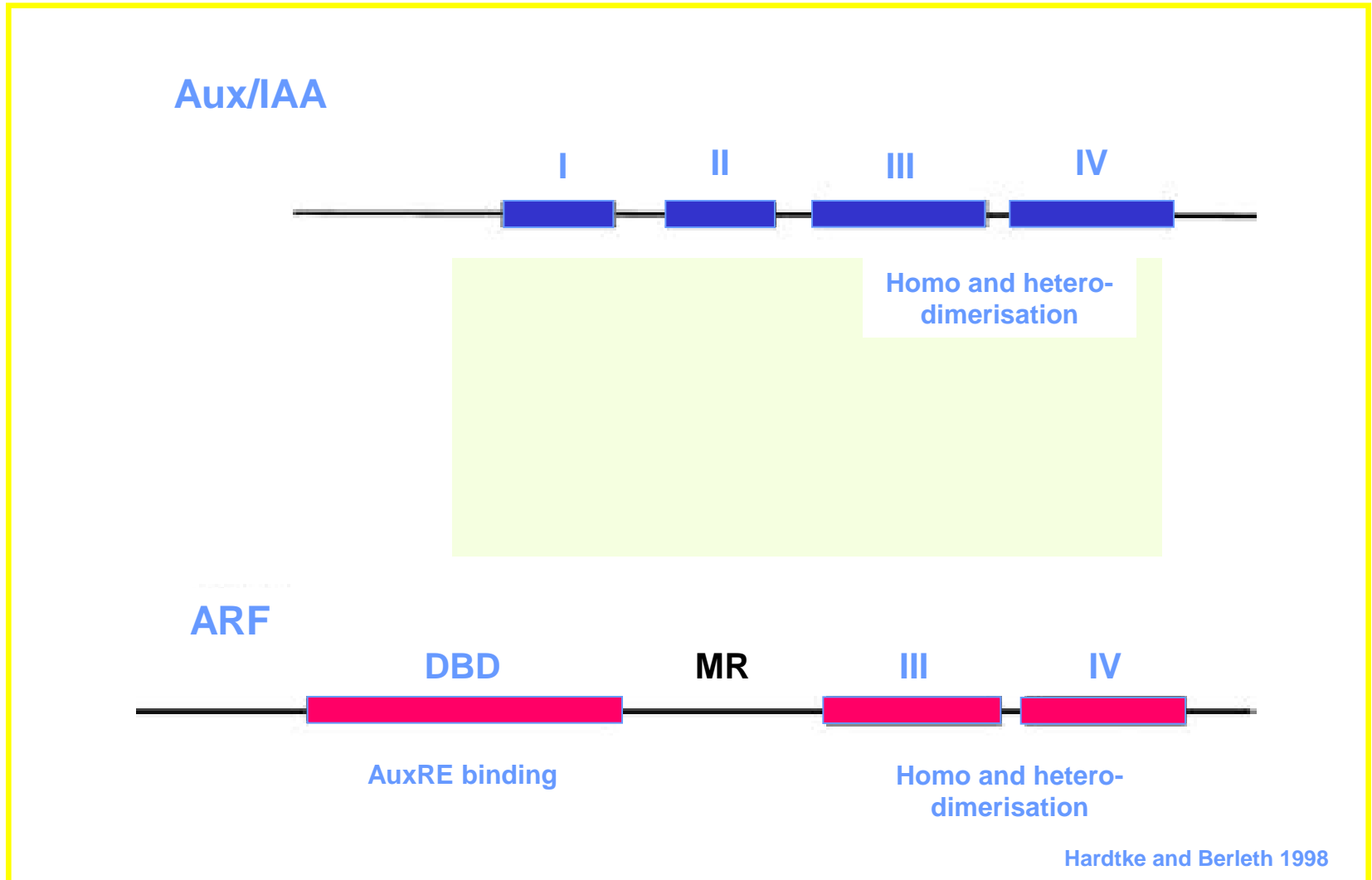
- Rapidly upregulated mRNAs
(*GH3*, *SAUR*, *AUX/IAA* genes)
- One hybrid screen with Auxin Response Elements
=> Auxin Response Factors (ARF)
- Two hybrid => *AUX/IAAs* interact with ARFs

Molecular Biology Approach to Elucidate Auxin Signaling

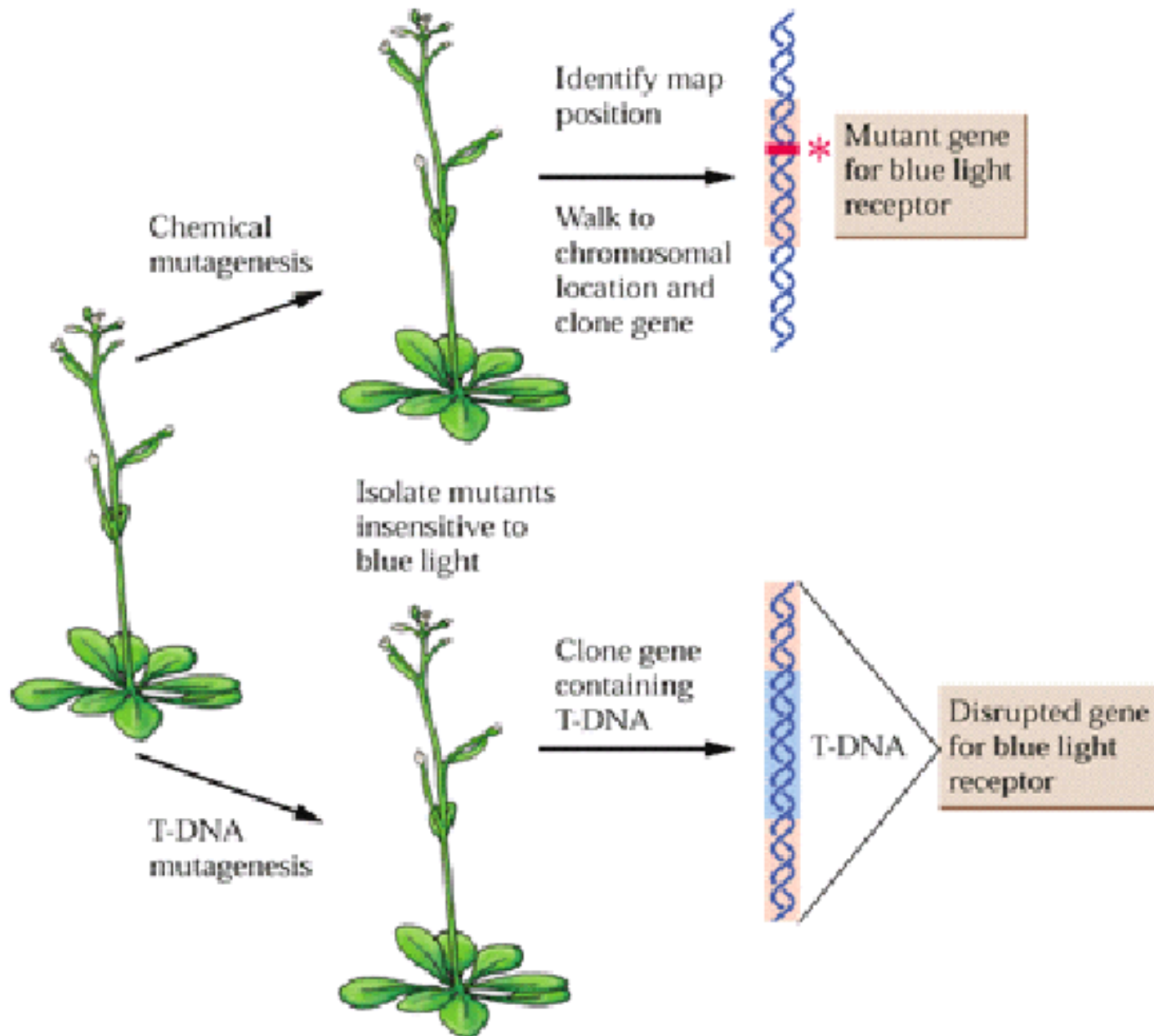
RT-PCR



ARFs auxin response factors – TF controlling, Aux/IAA expression



Forward genetics



Genetic Approach to Identify Auxin Receptor

- Auxin resistant (axr): *axr1* - *axr6*

- Transport inhibitor response (tir):
tir1 - *tir7*

Morphological mutants (*monopteros*, *bodenlos*,
etc.)

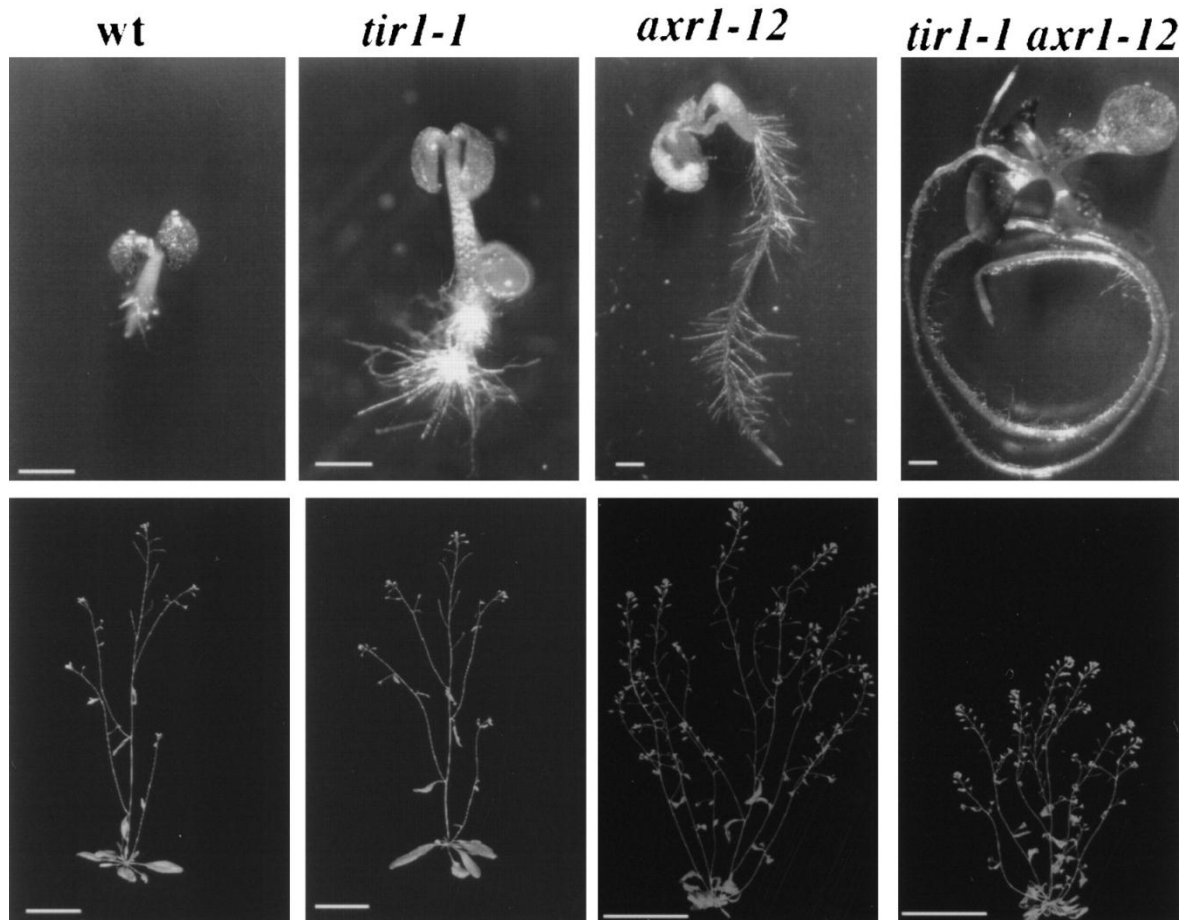
=> Role of regulated protein degradation and
transcriptional regulation in auxin
signaling

None of the identified gene looks like a
receptor

Auxin resistant (*axr*): *axr1* - *axr6*

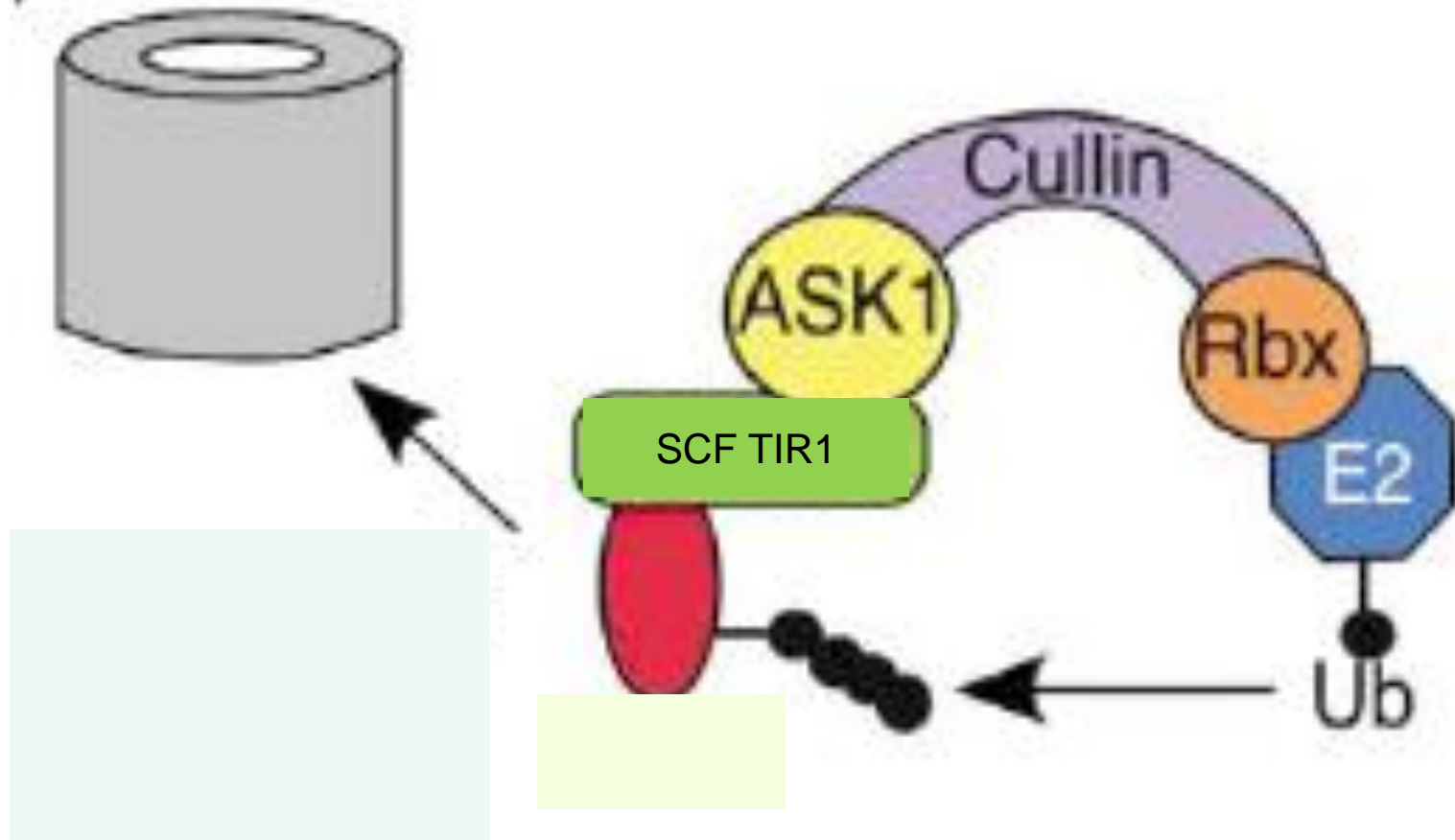
Auxin Transport inhibitor response (*tir*):

B

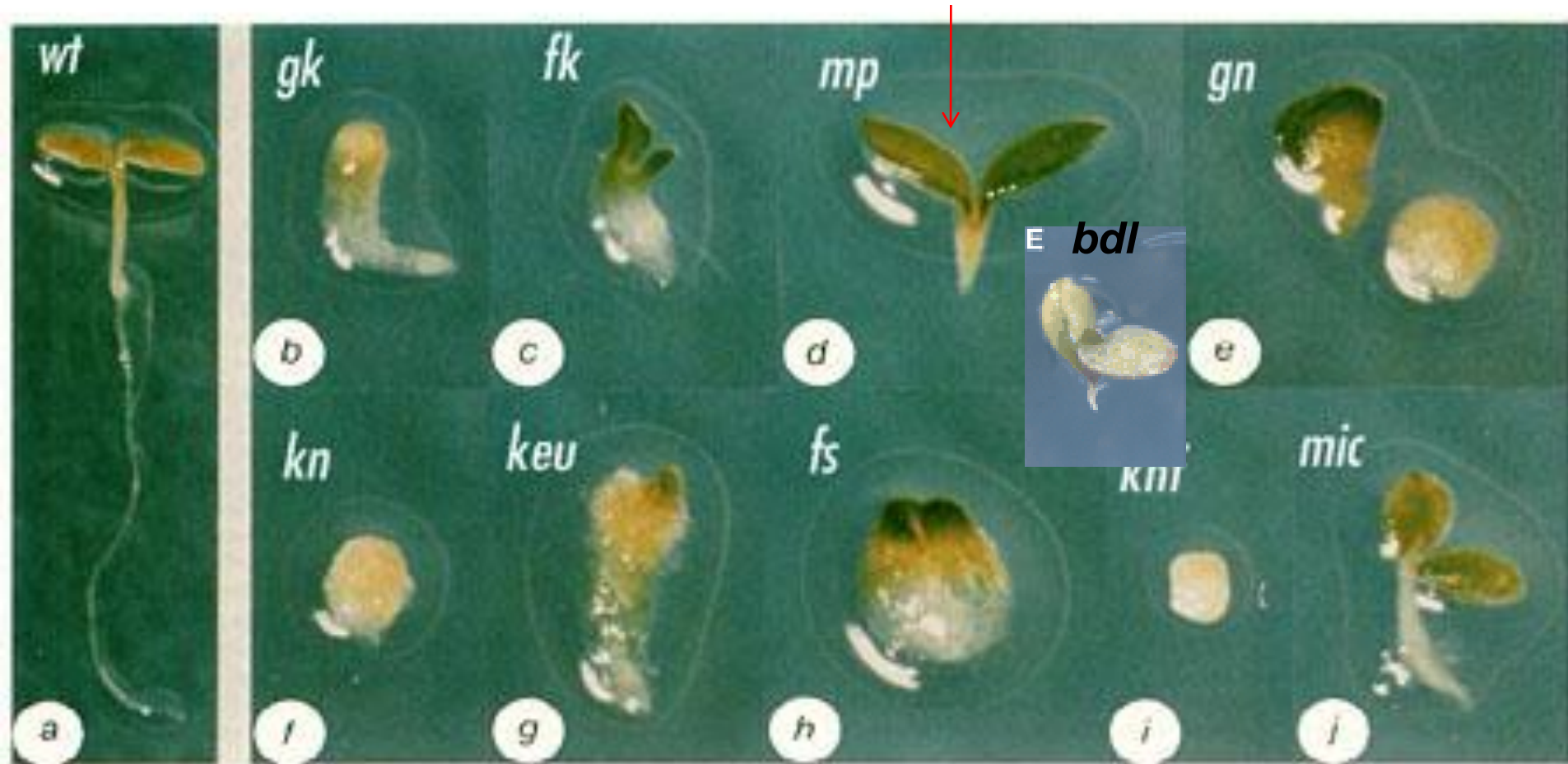


Subunits of ubiquitin ligase

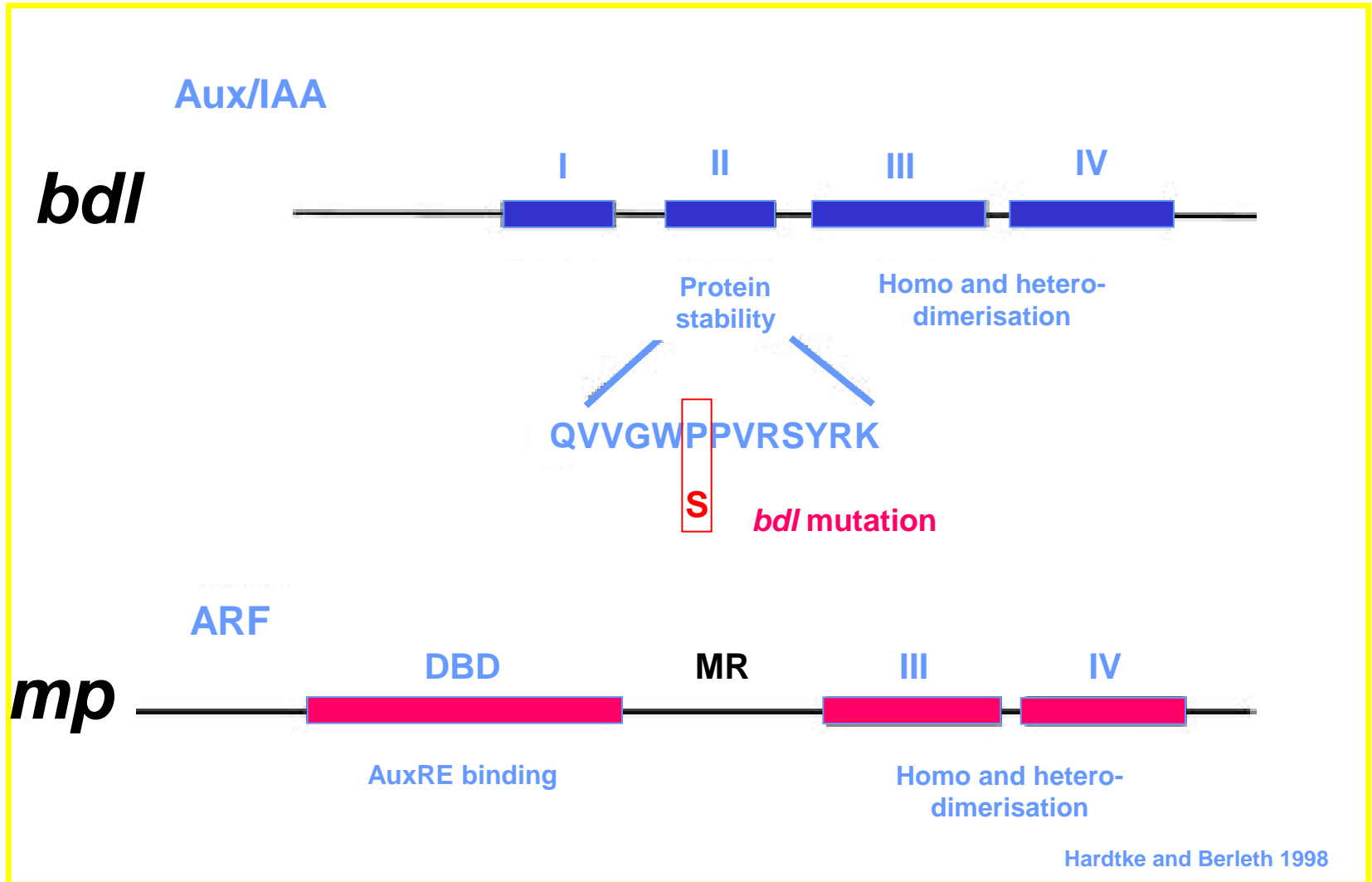
proteasome



Mutant Screen at Seedling Level



Some ARFs are **Activators**, and AUX/IAAs **Repressors** of Auxin Response



IAA13 *iaa13*^{P80S}

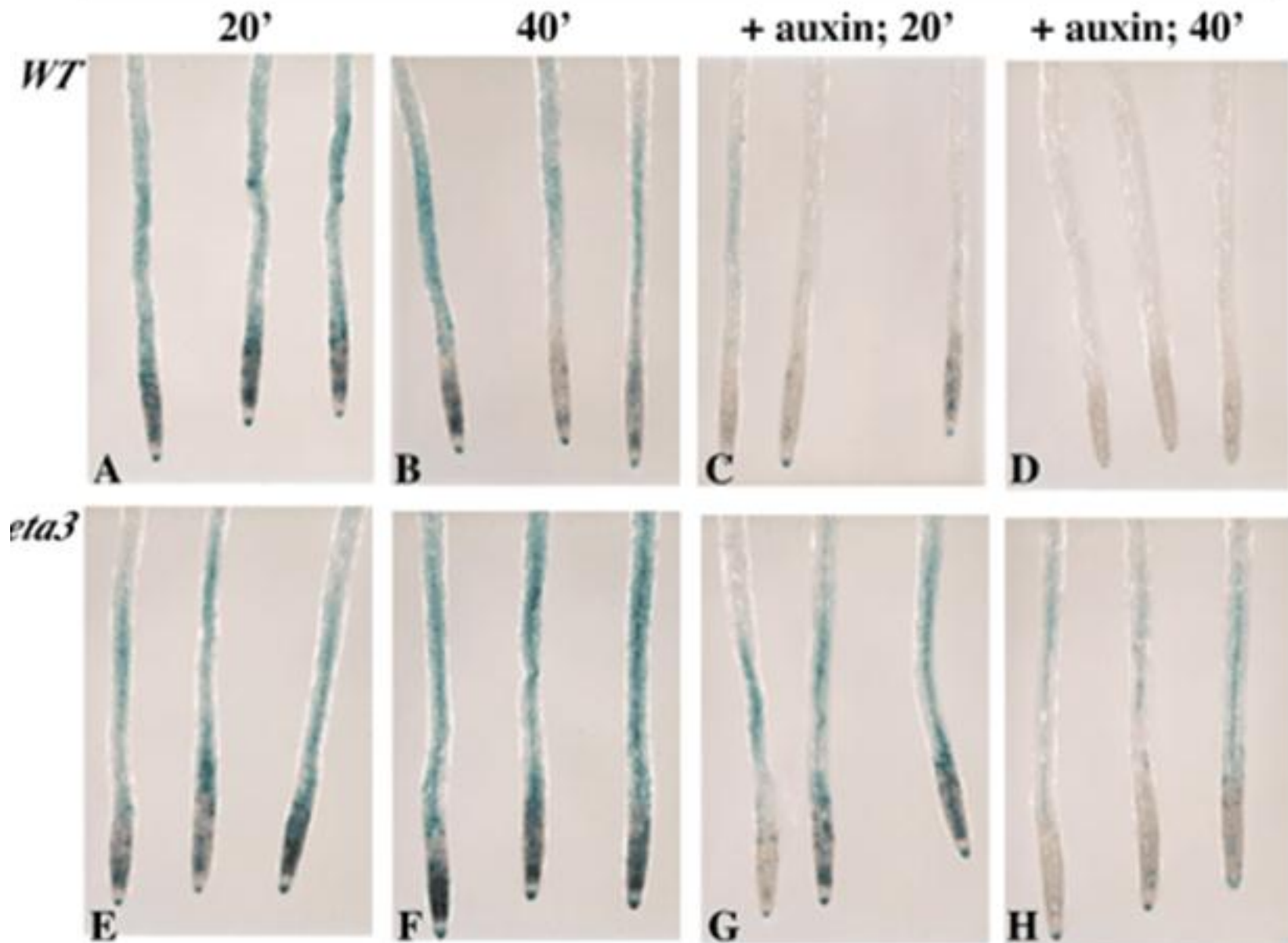


IAA13

*

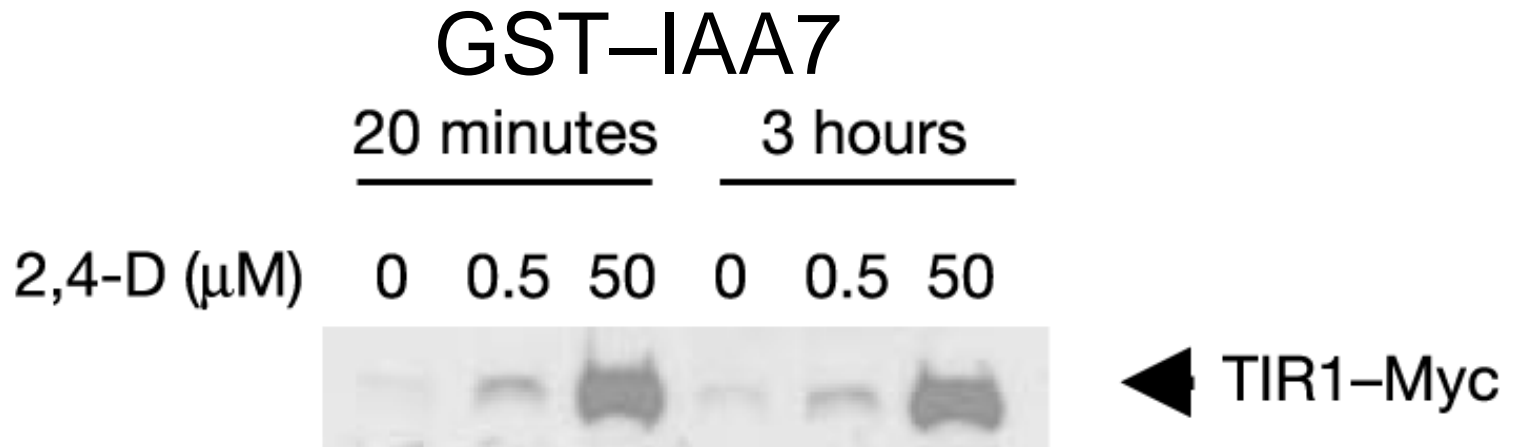
AUX/IAAs are stabilized in enhancer TIR1 mutant

HS:AXR3NT-GUS



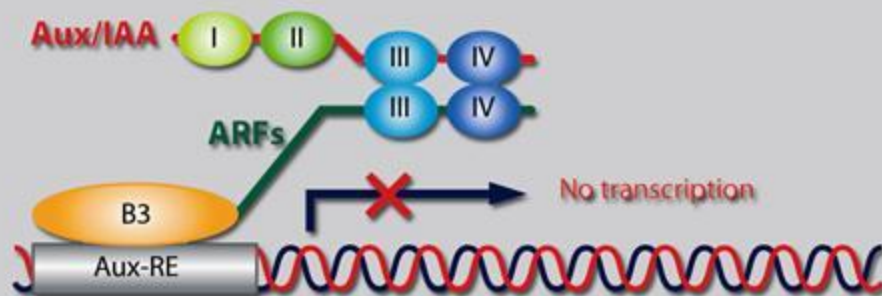
Pull-down

b

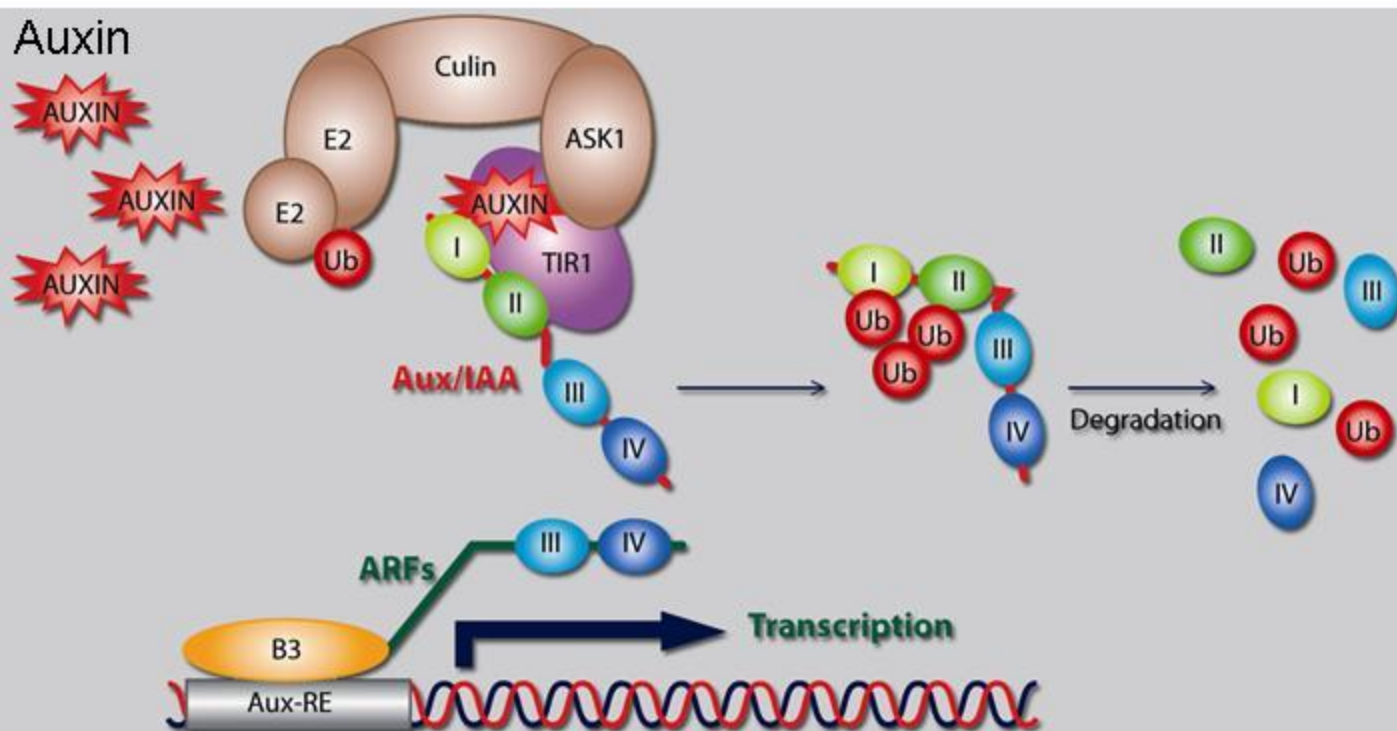


Auxin signaling

- Auxin



+ Auxin



Summary for Auxin Signaling

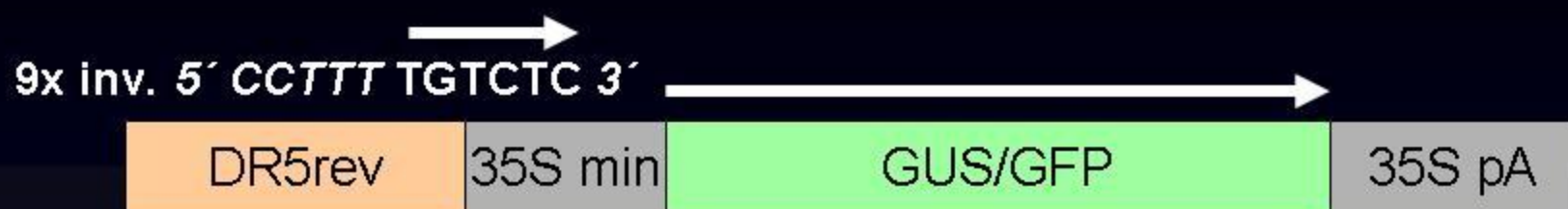
Biochemical approach - auxin binding protein
ABP1

binds auxin, important in embryogenesis,
role in endocytosis

Genetic approach - role of protein degradation
(*axr1*, *tir1*)

Molecular approach - auxin regulates expression
ARE in promoters 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



Root

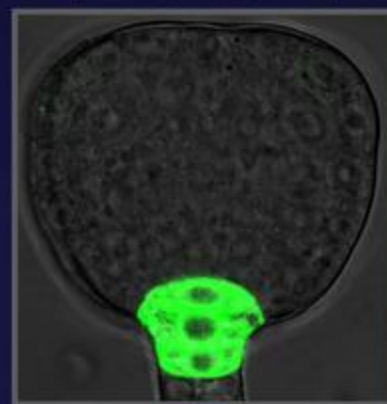
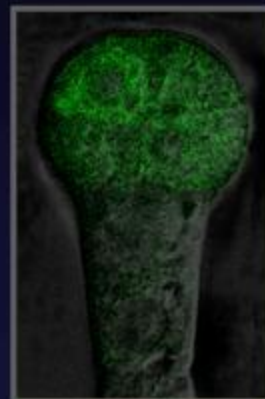
Embryos



DR5



anti-IAA

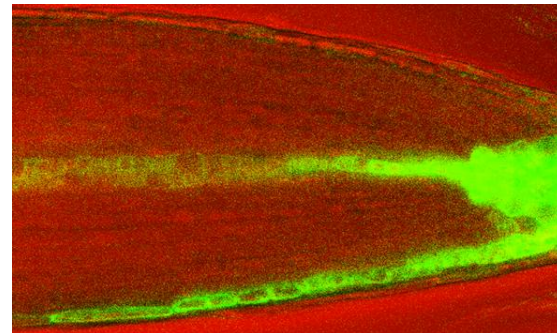
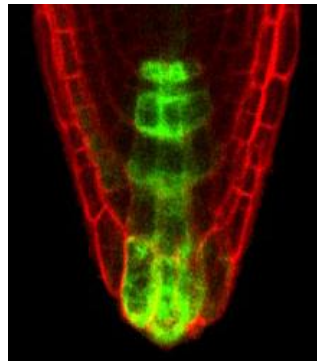
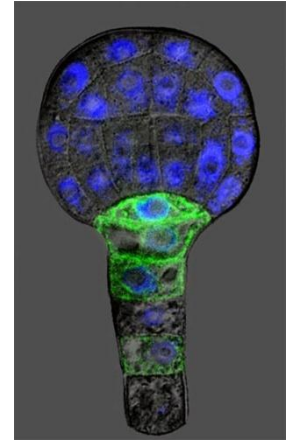
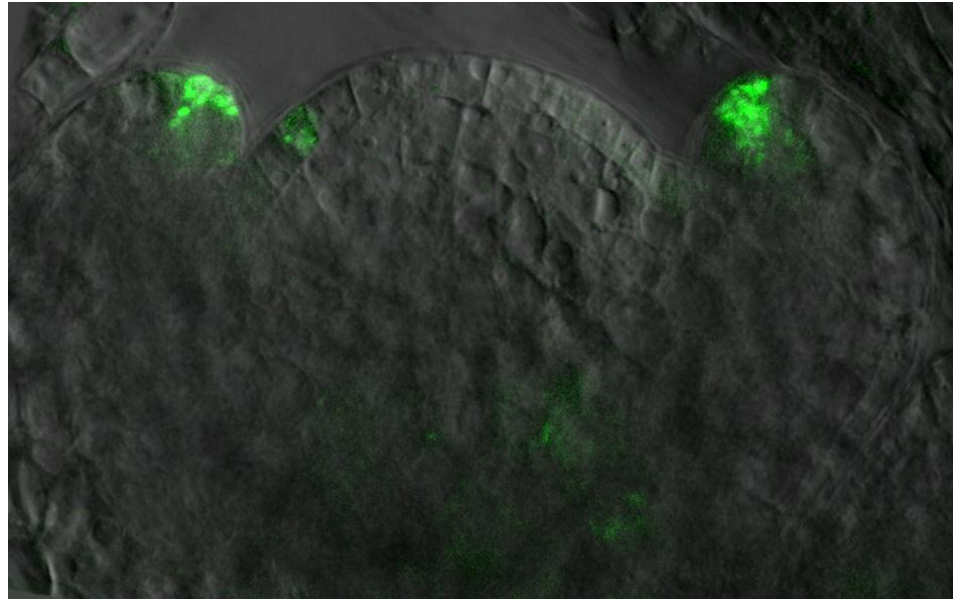
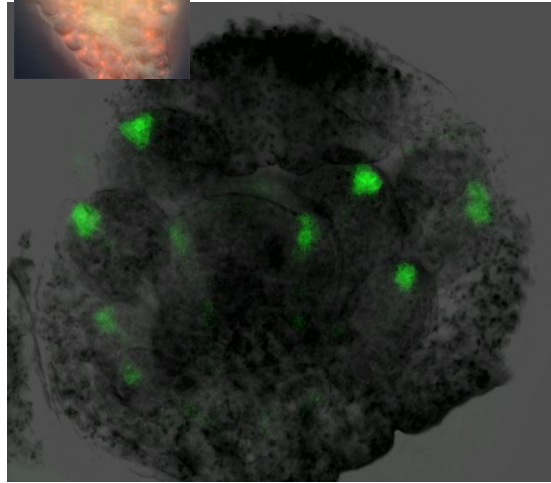


DR5

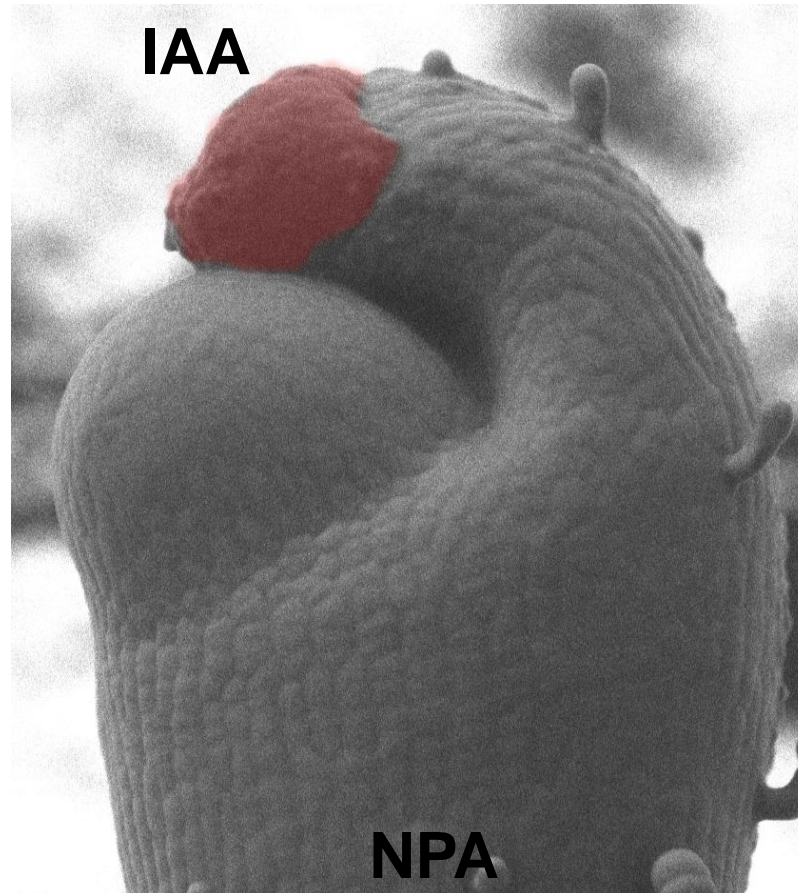
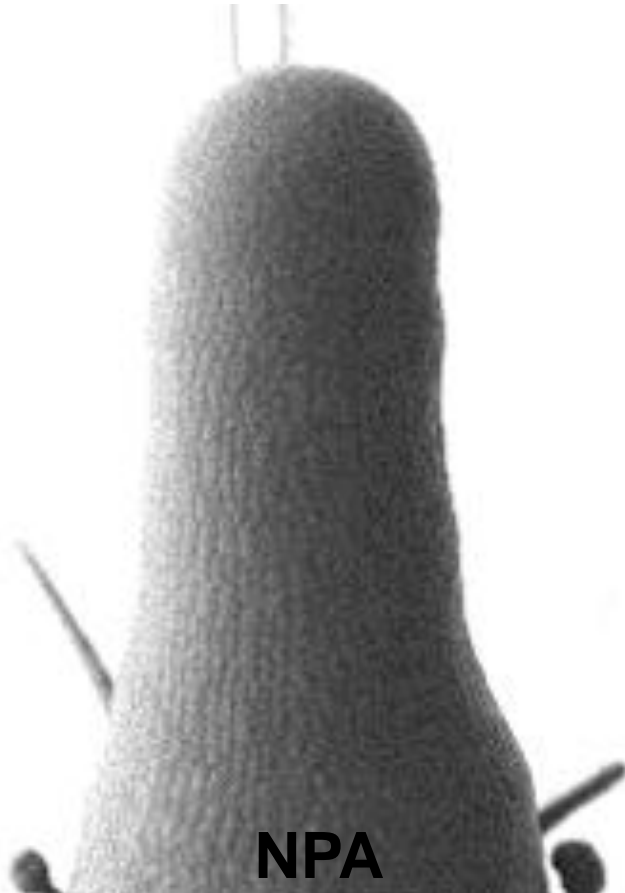


anti-IAA

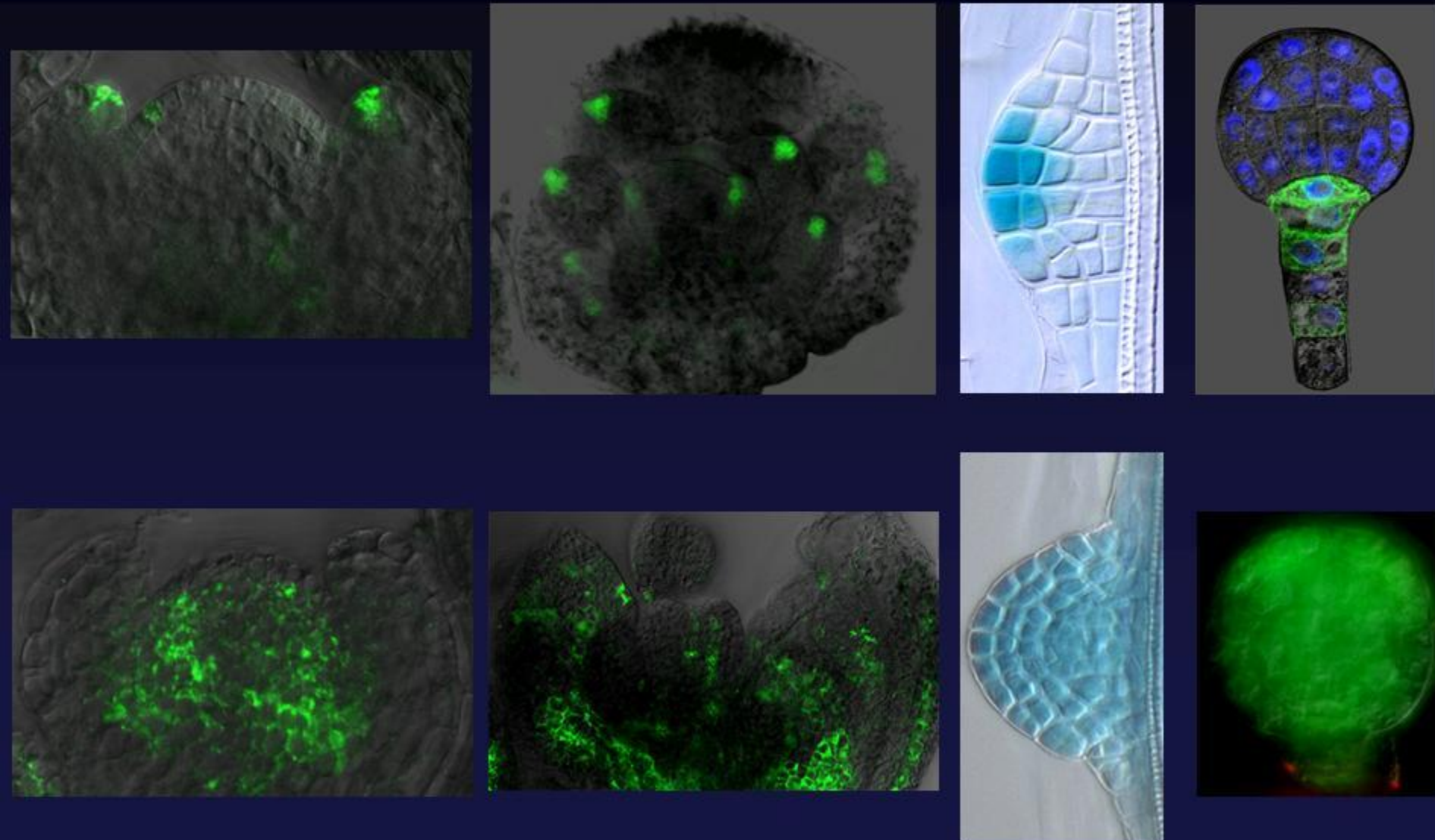
Local Auxin Gradients in Plant Development



Local Application of Auxin Induces Organ Formation



Local Auxin Gradients Require Active Polar Auxin Transport



Auxin Transport

Proteins involved in auxin transport

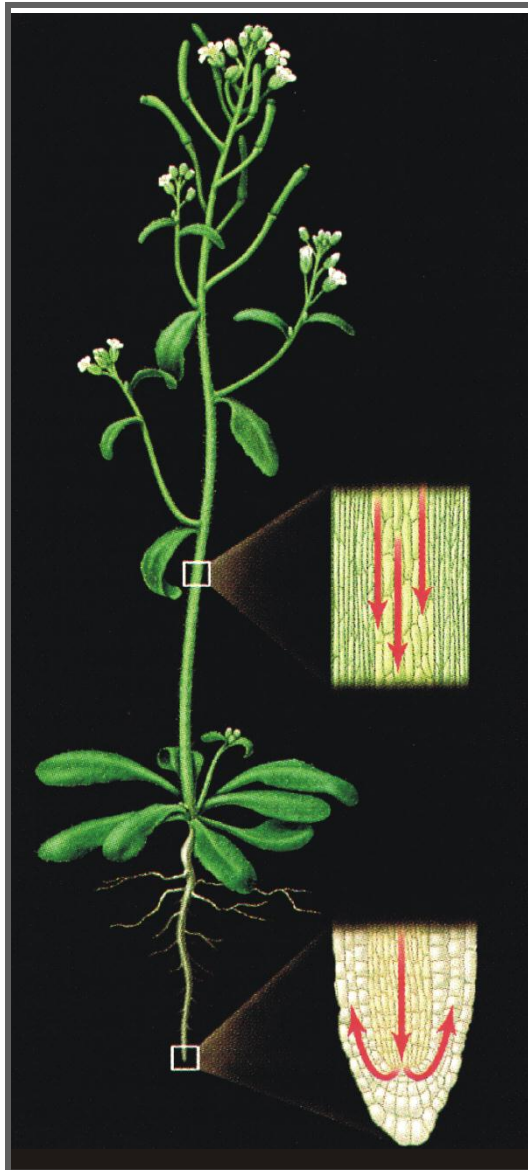
- PIN proteins (efflux)

- AUX1 proteins (influx)

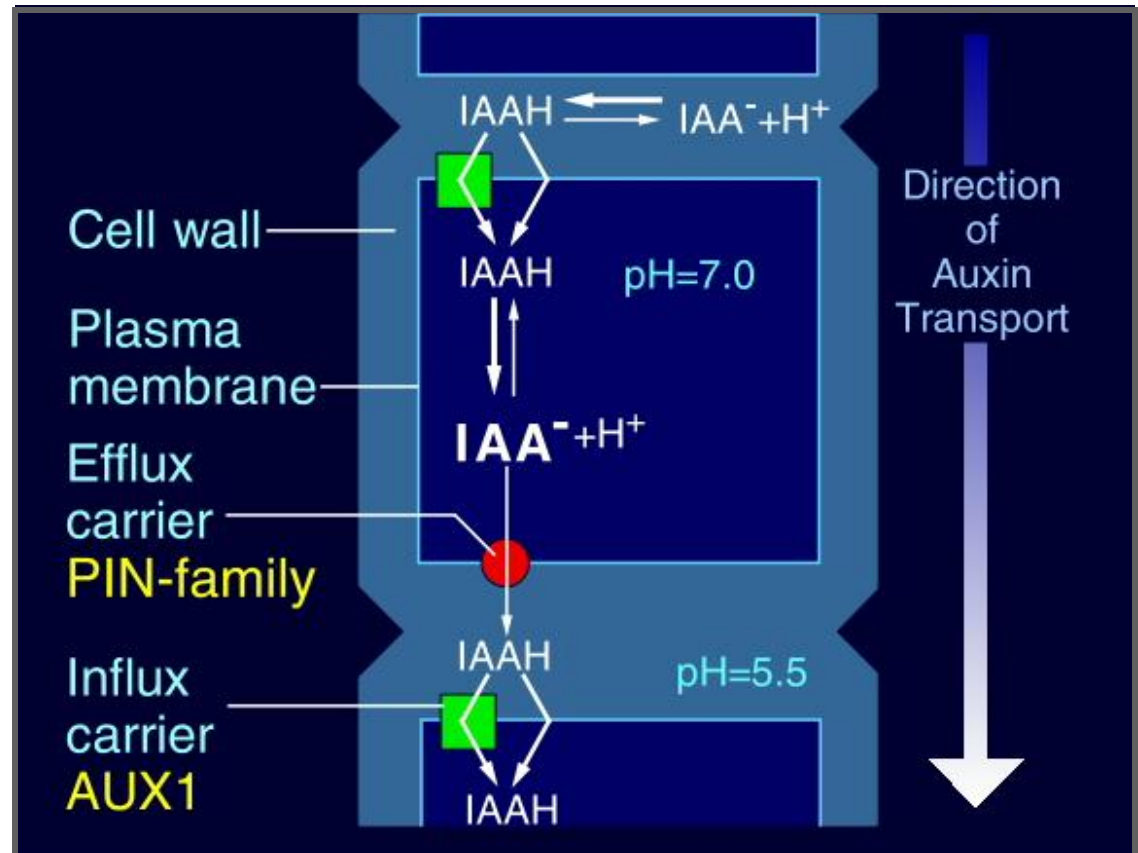
Role of *GNOM* dependent vesicle
trafficking

PIN proteins cycling and its role

Physiology of Auxin Transport



Chemiosmotic hypothesis

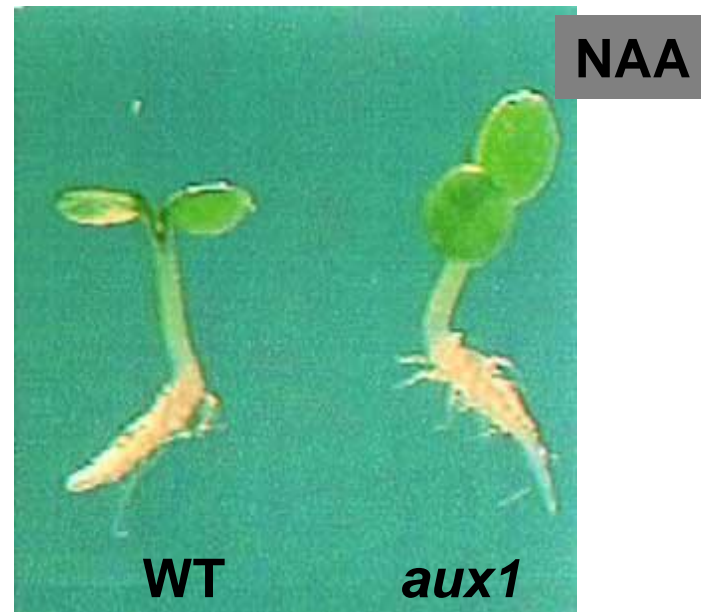
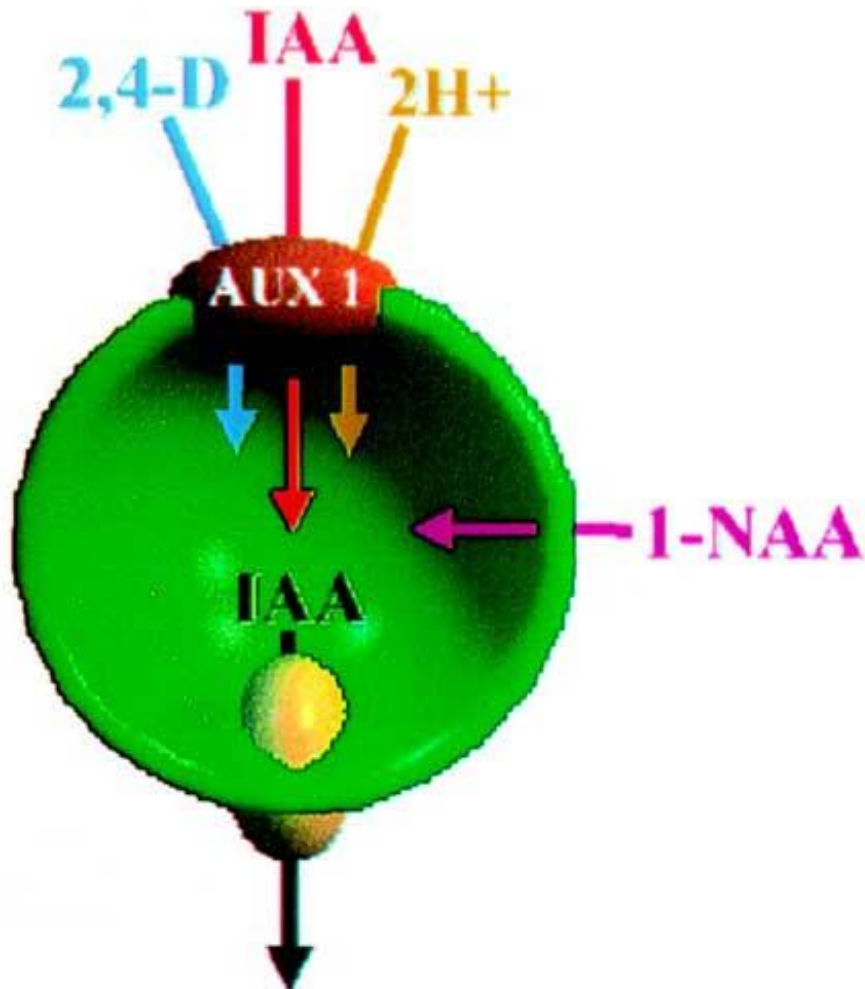


Auxin Influx

aux1 is Resistant to Auxin

aux1 phenotype

Transport properties of different auxins



NAA Rescues *aux1* Phenotype

- NAA



+ NAA



AUX1 – Expression and Localization

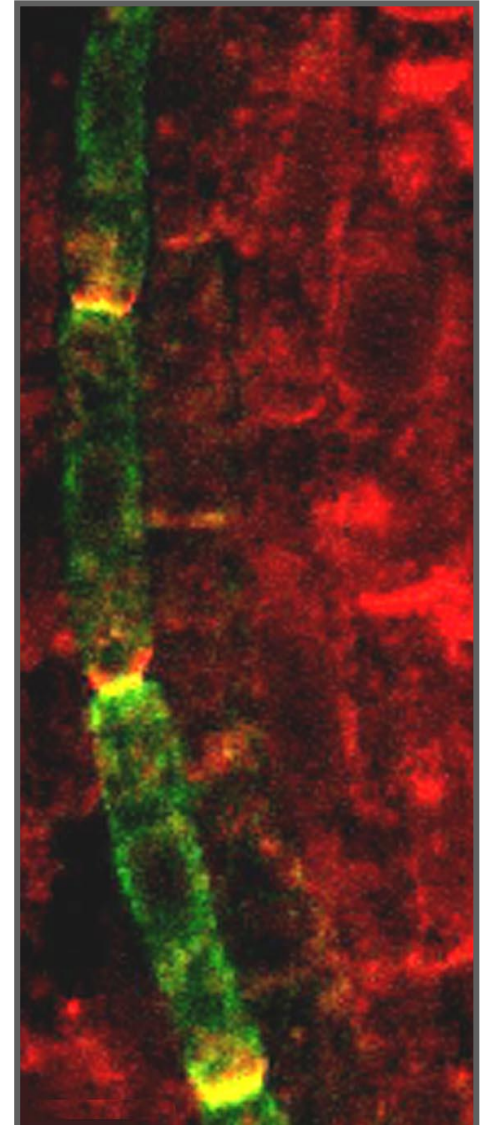
AUX1::GUS



AUX1 protein



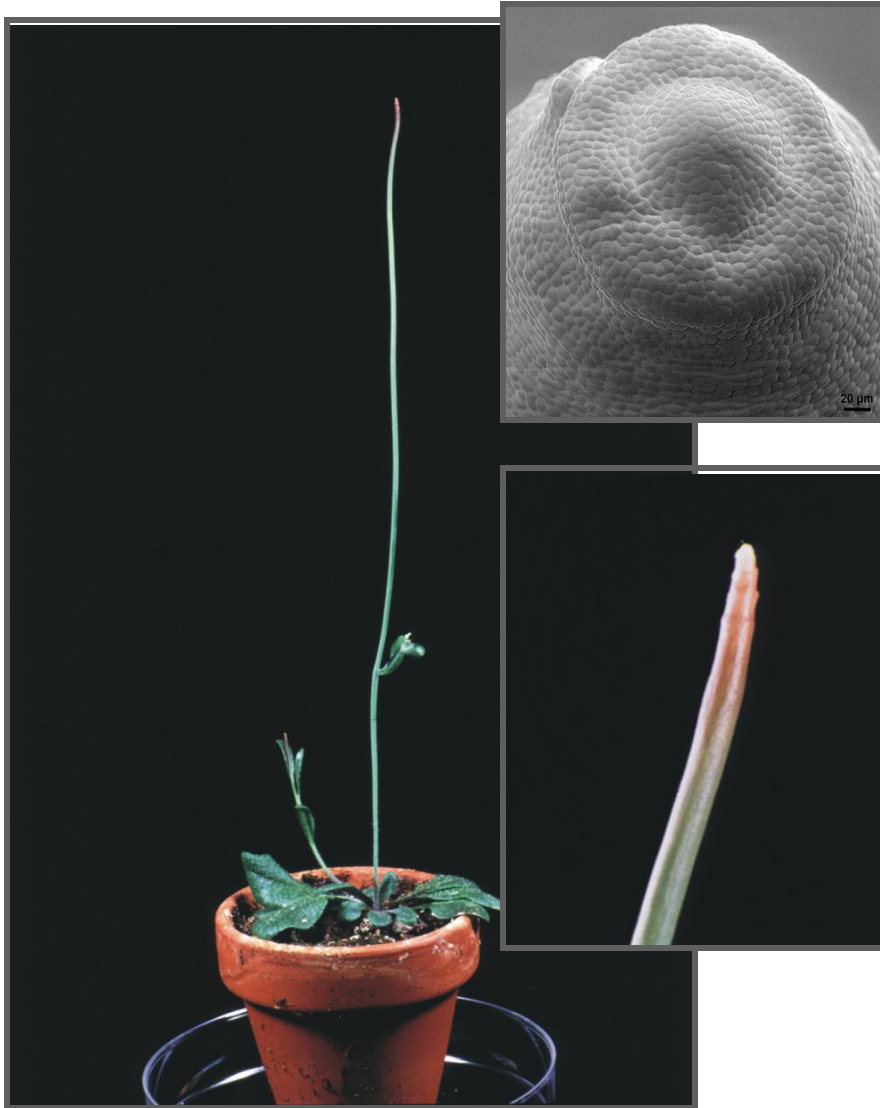
PIN1/AUX1



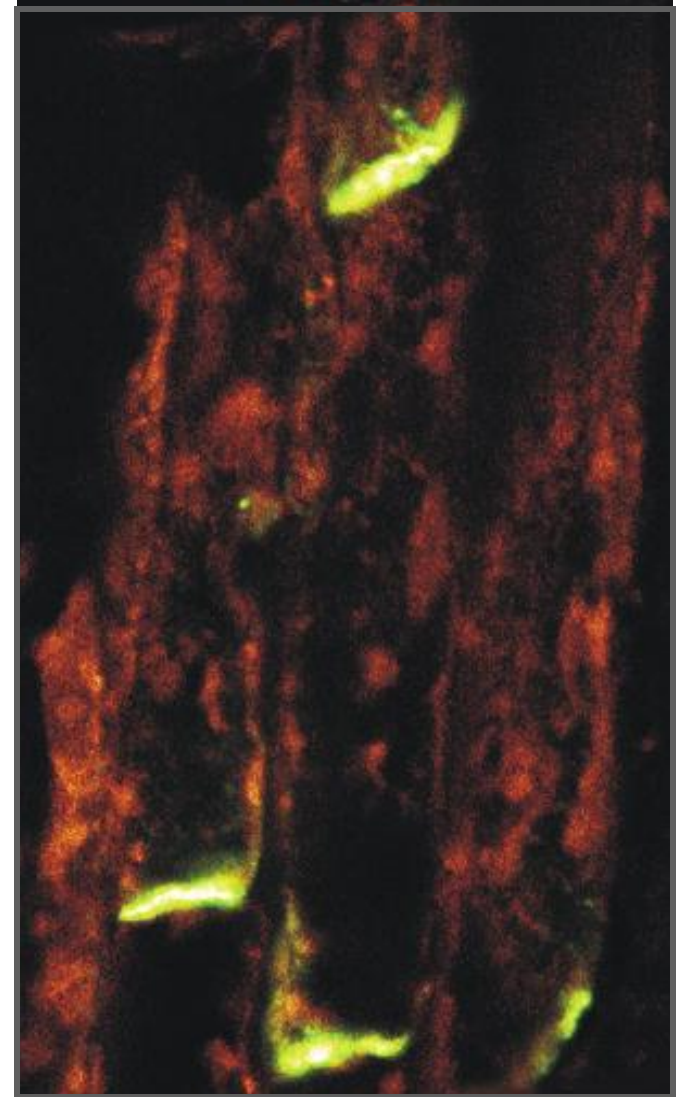
Auxin Efflux

PIN1 – the Auxin Efflux Carrier?

pin1 mutant

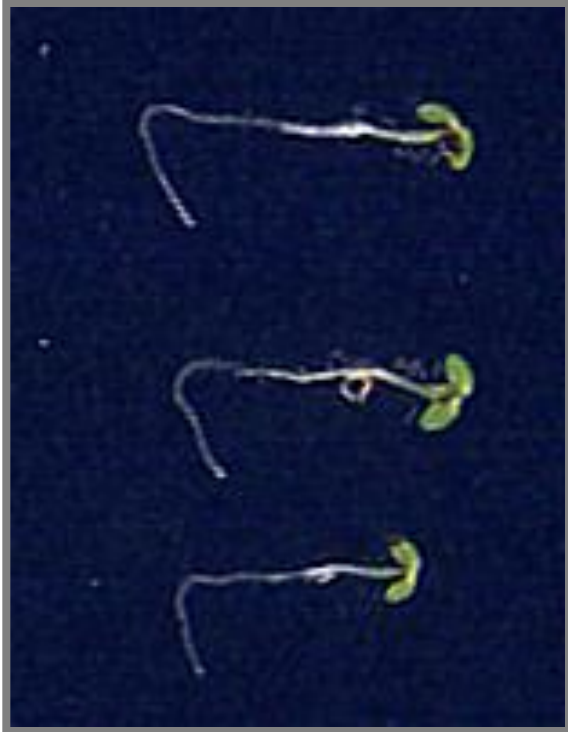


PIN1 protein



PIN2 – Root Gravotropism

Col-0



pin2



PIN2 protein

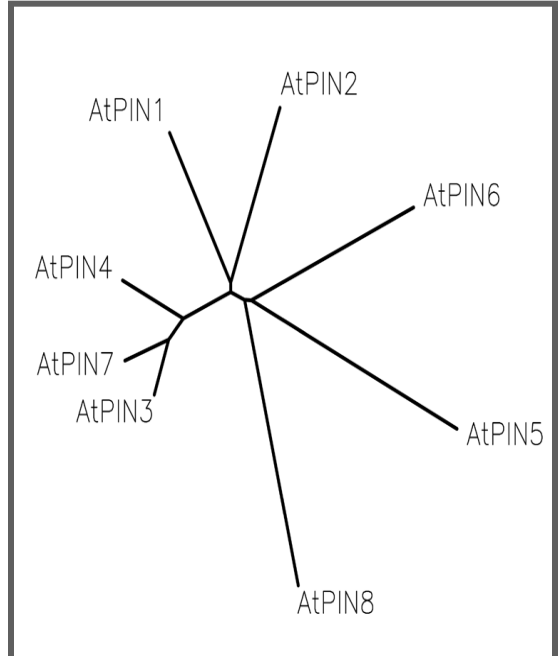


The Arabidopsis PIN Gene Family

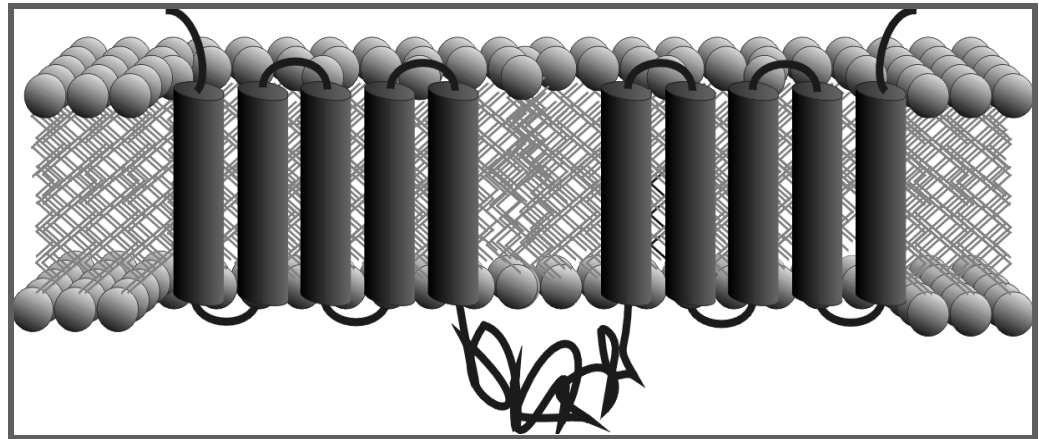
Comparison of Arabidopsis PIN proteins

| | | |
|--------|---|---|
| AtPIN1 | 1 | MISAASTFVHMEAMVFLVAMILAYGSKWKEIFPDQCSGIRRFVALFAVPLLSFHFPAANNFYAMHLFEAADGQRVYVSESSP...ECKLRSNRS...EDWTFELFSLSELPNTLY |
| AtPIN2 | 1 | MISGKDMYDVAAMVELVAMILAYGSRVWKKIETPDQCSGIRRFVAVFAPVLLSFHFISNDPYAMMYHFLAADLQKLVVILAAEF...GOANSSRRGS...LEWMTLFLSLSLPTNTLY |
| AtPIN3 | 1 | MISRHDLVTVLTAIVPLVAMILAYGSRVWKKIETPDQCSGIRRFVAVFAPVLLSFHFISTNPNYAMHLRFIAADLQKLIIMLSLVL...WANFTRSSG...LEWMTLFLSLSLPTNTLY |
| AtPIN4 | 1 | MISRHDLVTVLTAIVPLVAMILAYGSRVWKKIETPDQCSGIRRFVAVFAPVLLSFHFISTNPNYAMHFRFAADLQKLIIMLVLEAL...WANLTKRNS...LEWMTLFLSLSLPTNTLY |
| AtPIN5 | 1 | MISGCVYVYLIAMKELVAVLQVGSYKWKIEETPDQCSGIRRFVAVFAPVLLSFHFISTNPNYAMHFRFAADLQKLIIMLVLEAL...WANLTKRNS...LEWMTLFLSLSLPTNTLY |
| AtPIN6 | 1 | MISGNEPYSVCMACALIFAMFVAYGSKWKIKETPDQCSGIRRFVAVFAPVLLSFHFISQNNNYKMDTFMLADLTKIFVFLVLS...NAVFFKAGG...IDMLTLFLSLSLPTNTLY |
| AtPIN7 | 1 | MISRHDLVTVLTAIVPLVAMILAYGSRVWKKIETPDQCSGIRRFVAVFAPVLLSFHFISQNNNYKMDTFMLADLTKIFVFLVLS...NAVFFKAGG...LEWMTLFLSLSLPTNTLY |
| AtPIN8 | 1 | MISRHDLVTVLTAIVPLVAMILAYGSRVWKKIETPDQCSGIRRFVAVFAPVLLSFHFISQNNNYKMDTFMLADLTKIFVFLVLS...NAVFFKAGG...LEWMTLFLSLSLPTNTLY |

Phylogenetic tree



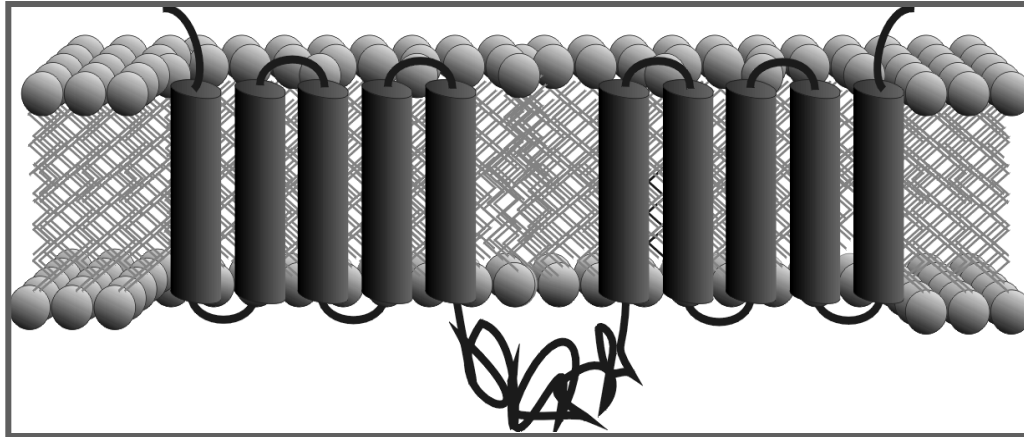
Membrane topology model



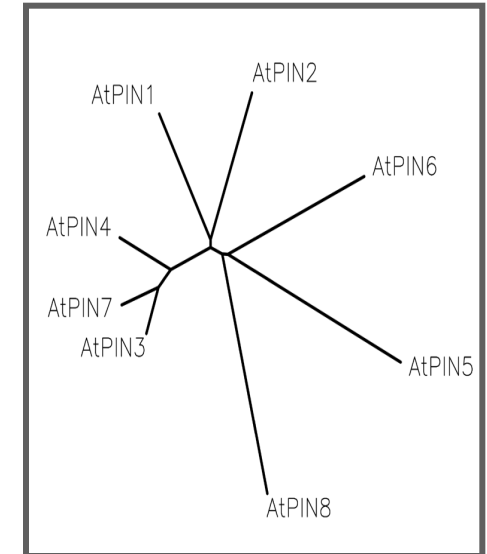
What is Molecular Role
of PIN Proteins
in Auxin Transport?

PINs Are Essential Components of Auxin Transport

Putative topology of PIN proteins



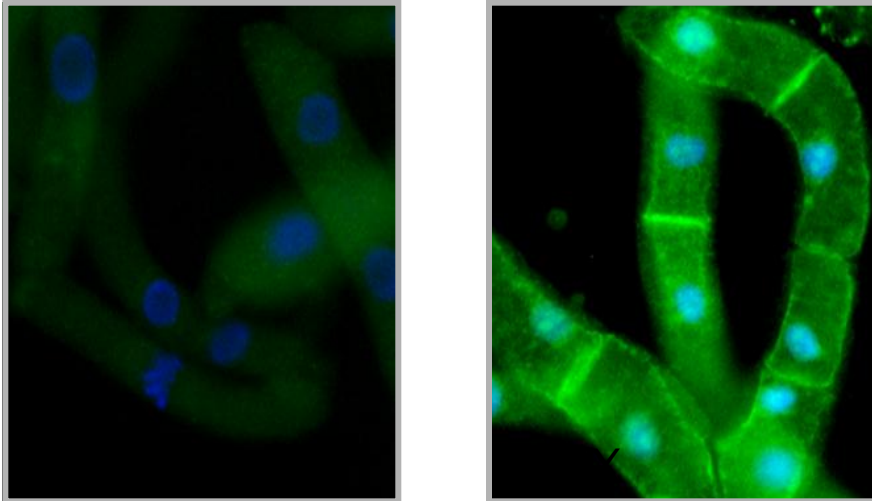
Phylogenetic tree



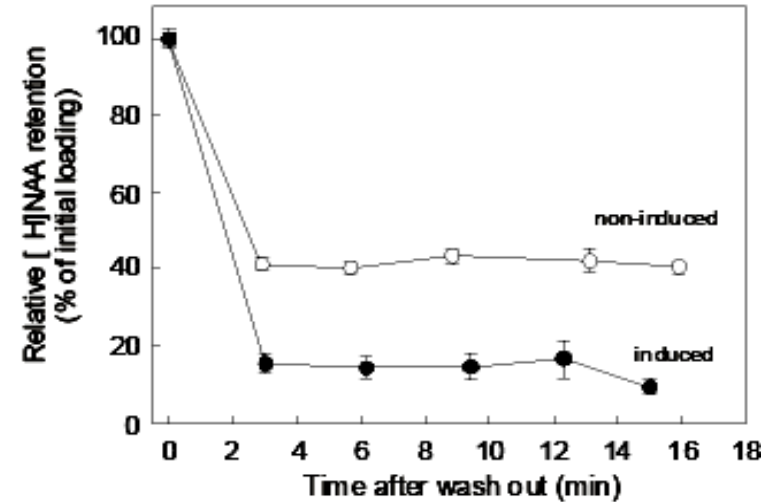
- All defects in *pin* loss-of-function mutants are in auxin transport-dependent processes and can be phenocopied by auxin transport inhibitors
- Local auxin distribution (gradients) are affected in *pins*
- Polar PIN localization determines direction of auxin flow

PINs Are Rate-limiting Factors in Auxin Efflux

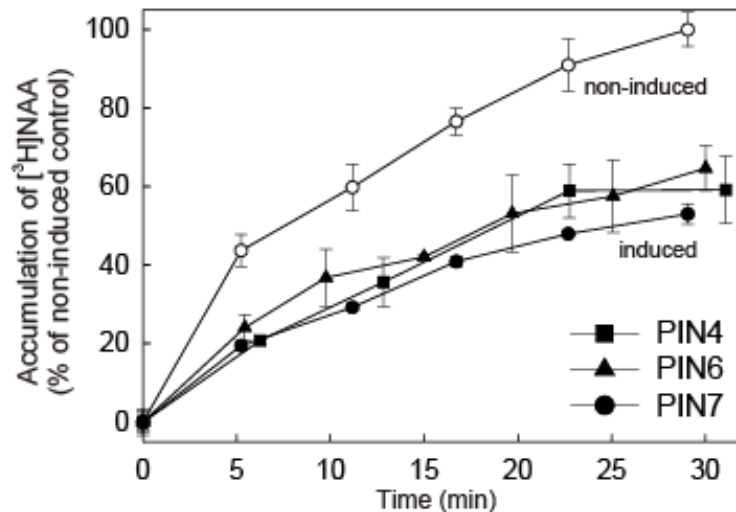
Inducible PIN1 expression



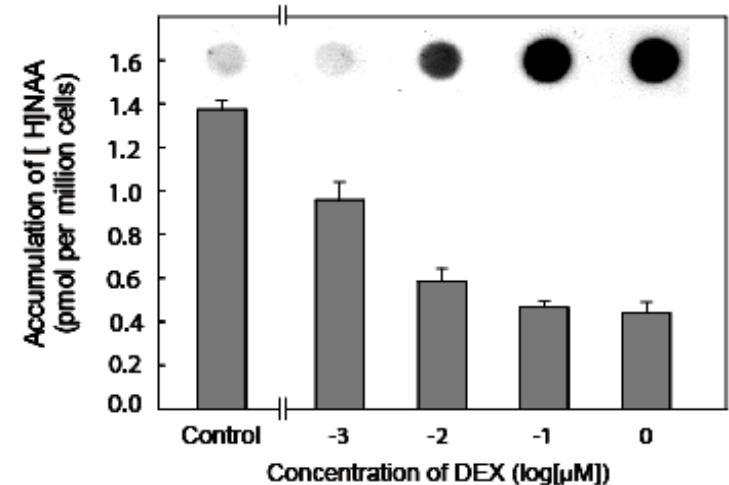
PIN-dependent auxin efflux from GVG-PIN7 tobacco cells



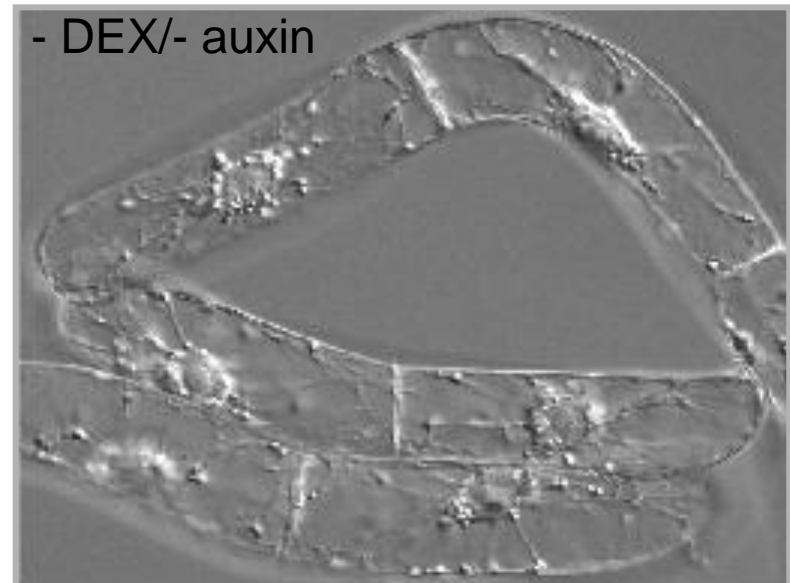
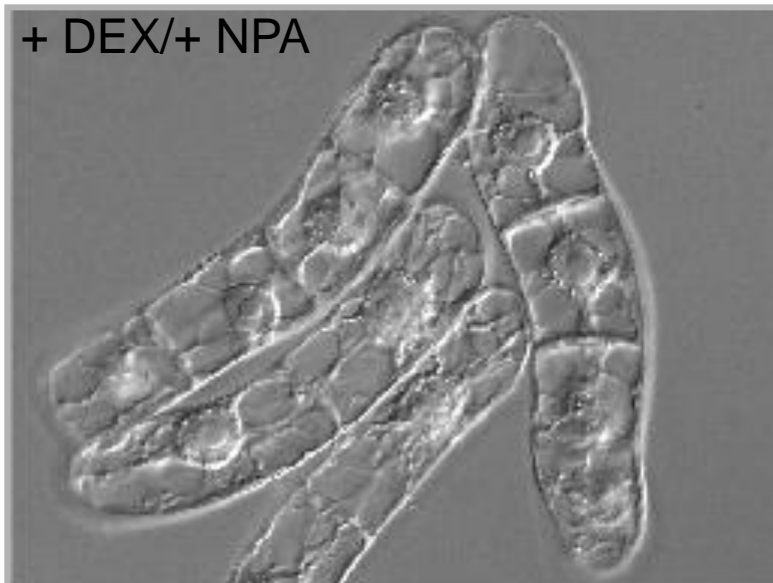
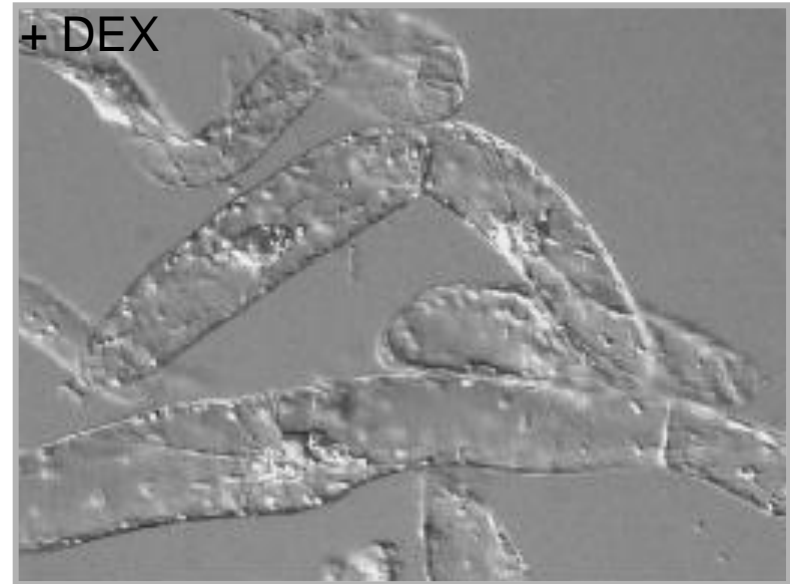
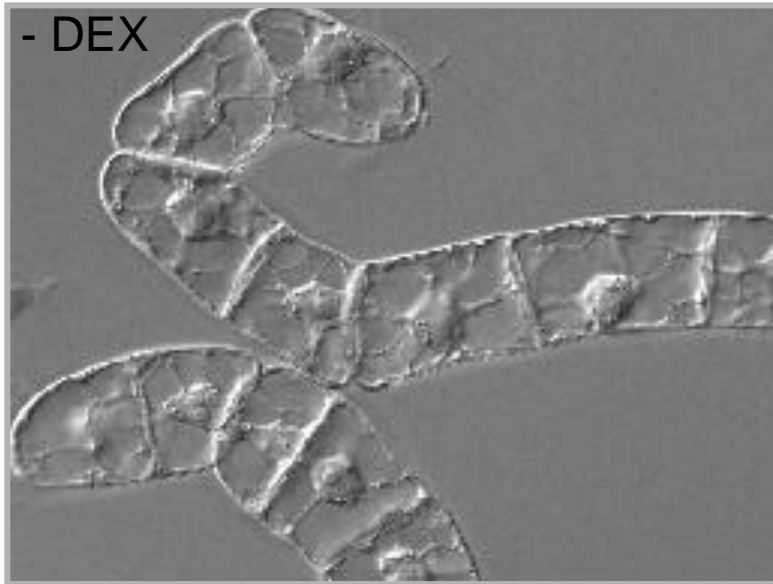
NAA accumulation kinetics



[3H]NAA accumulation in GVG-PIN7 tobacco cells in relation to DEX concentration

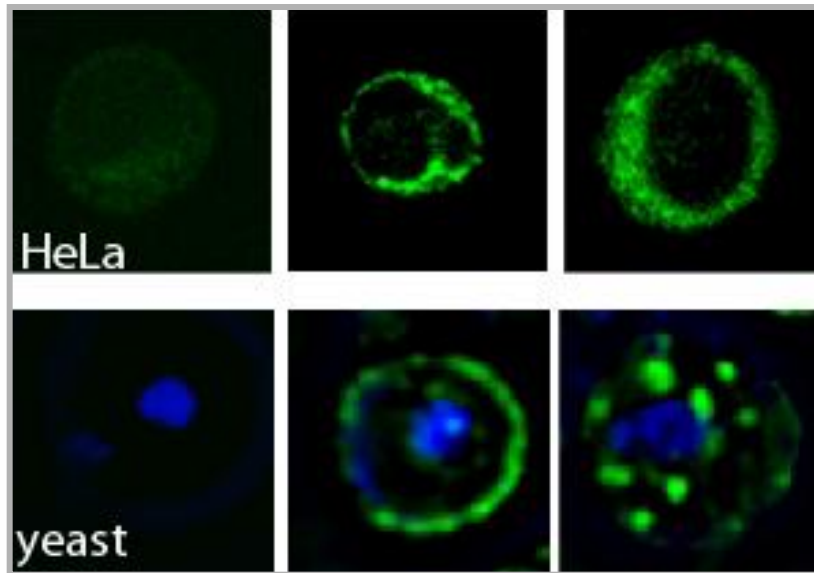


PIN-induced Phenotypes in BY-2 Cells



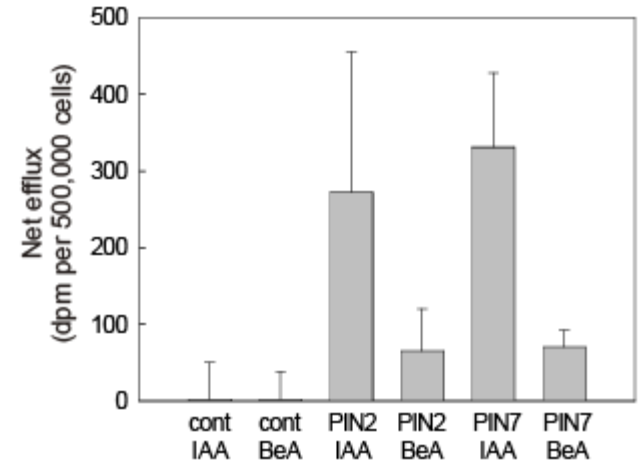
Expression of PINs in HeLa and Yeast

Heterologous PIN2 expression

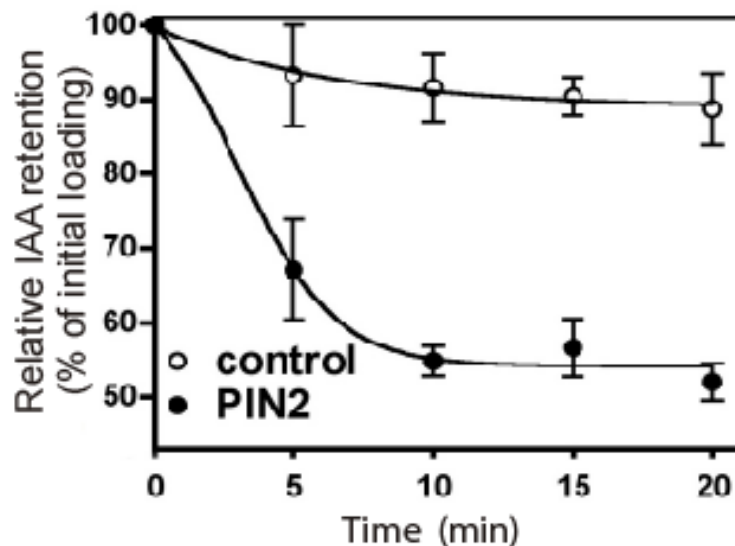


auxin efflux activity

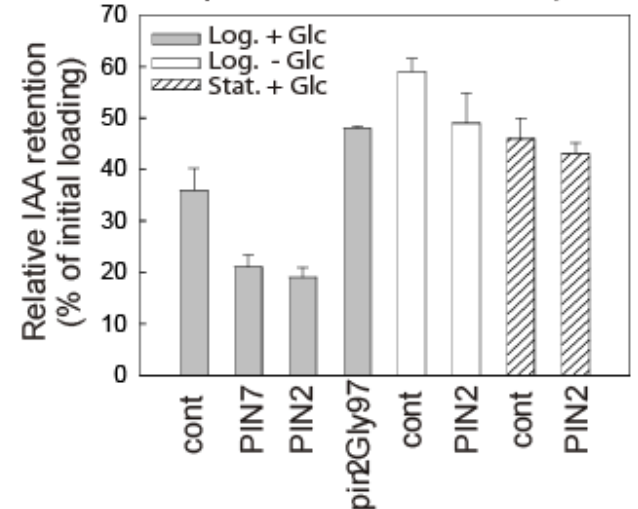
PIN-dependent efflux in HeLa cells



IAA ef flux in yeast



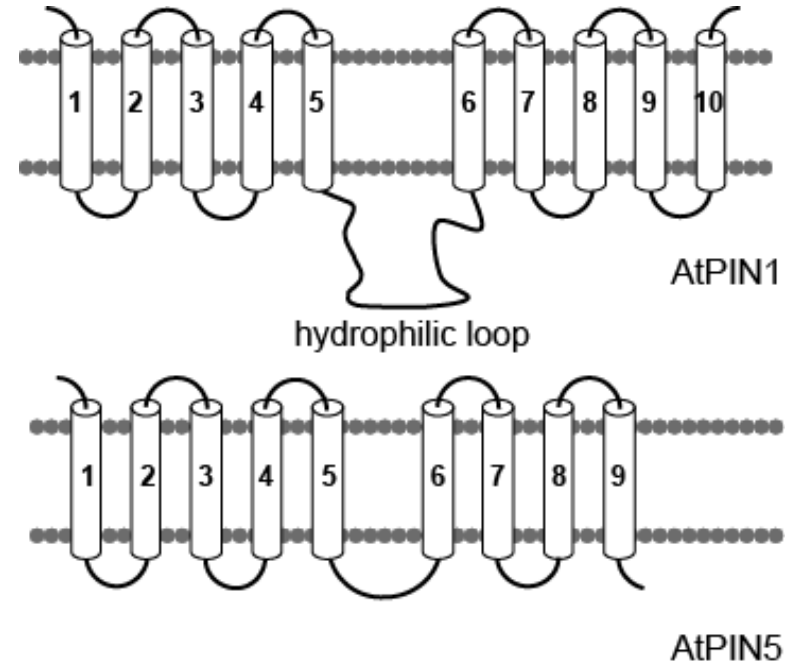
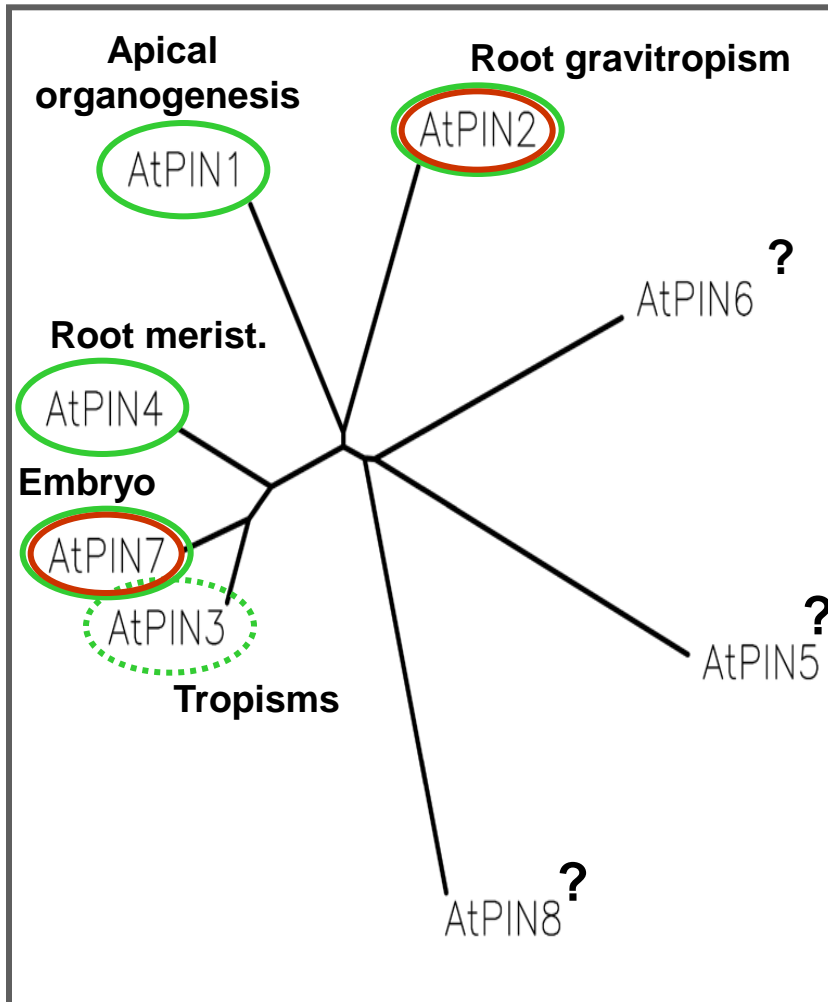
PIN-dependent IAA ef flux in yeast



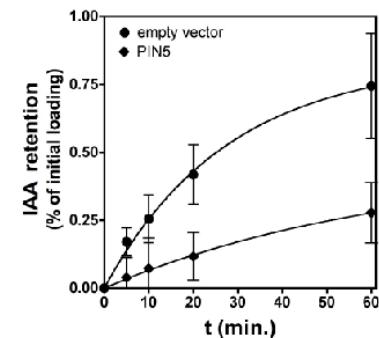
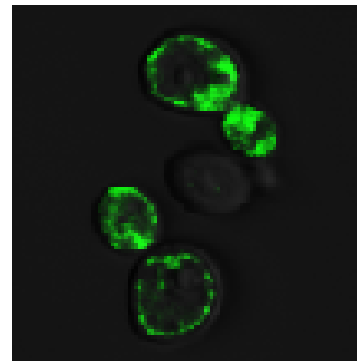
PIN gene family

Predicted PIN Protein Topology

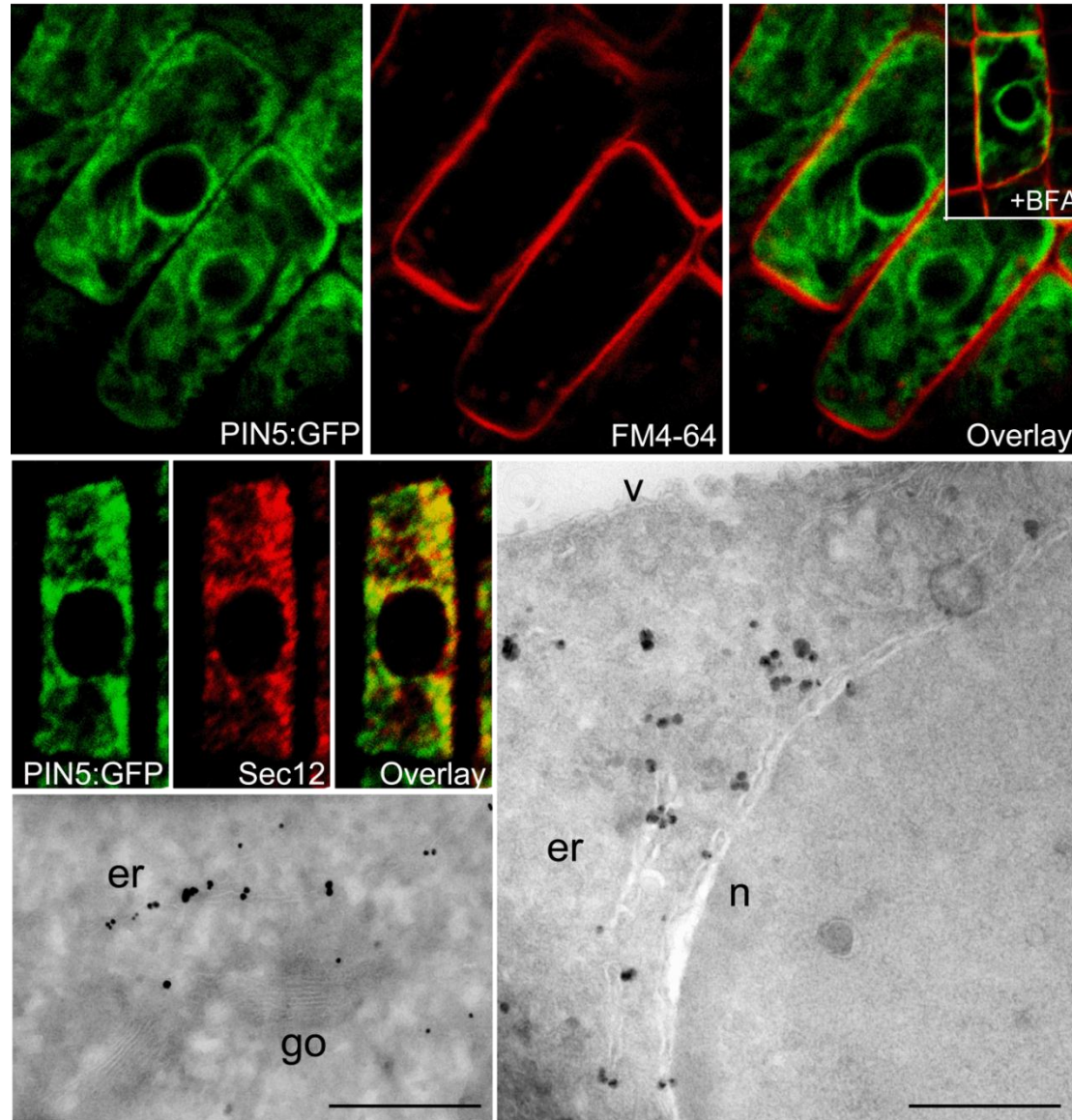
PINs in *Arabidopsis*



Auxin Transport in Yeast

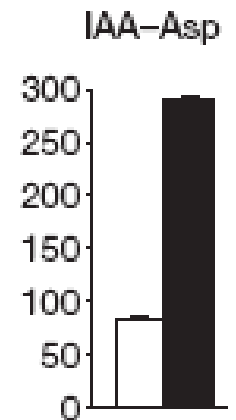
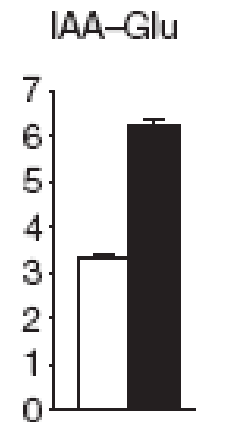
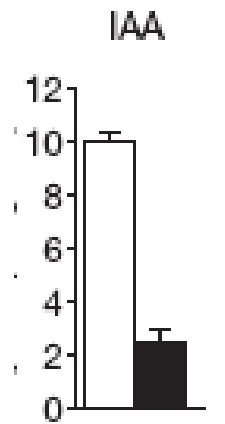
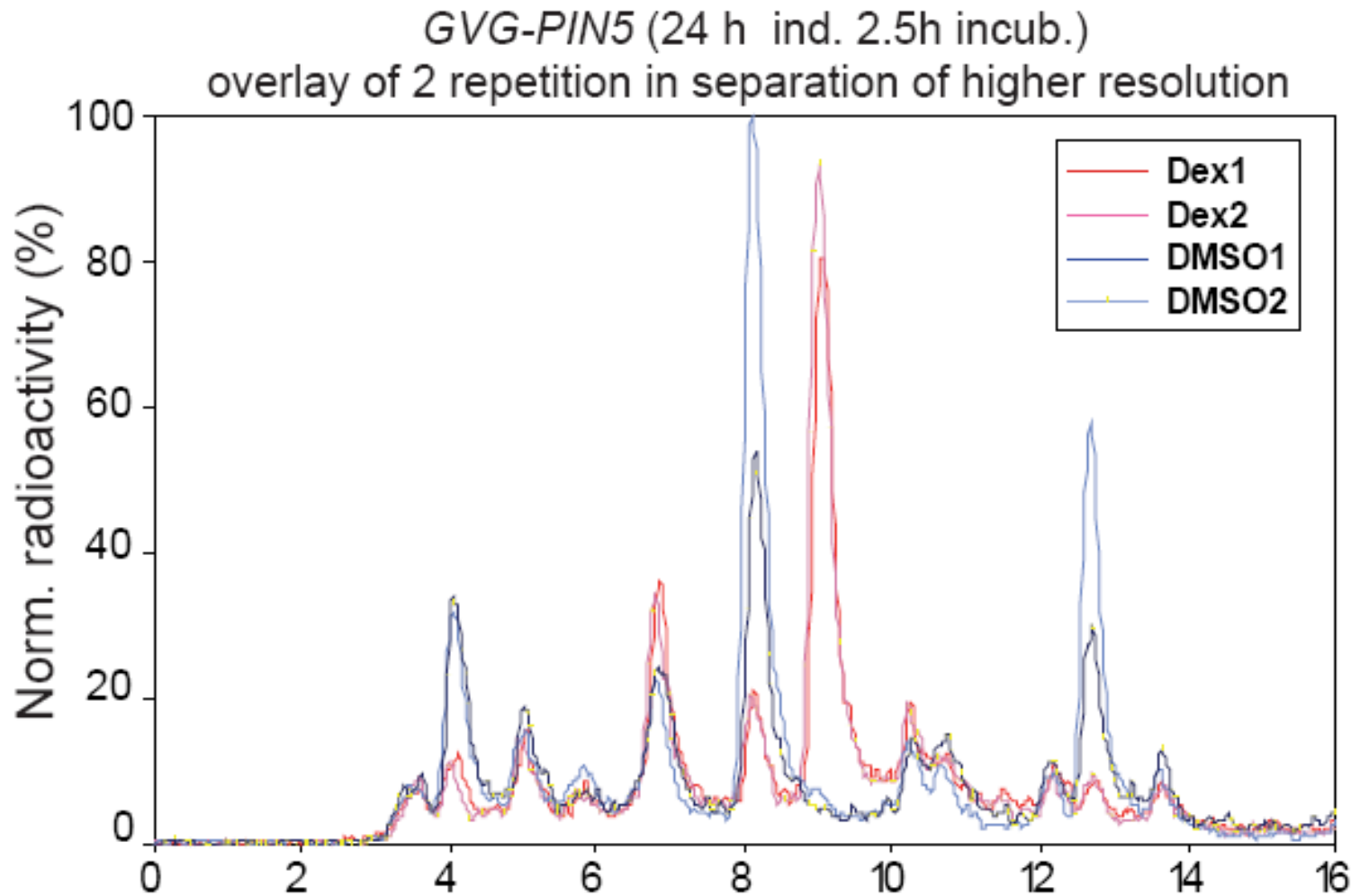


ER-based PIN5-dependent auxin transport

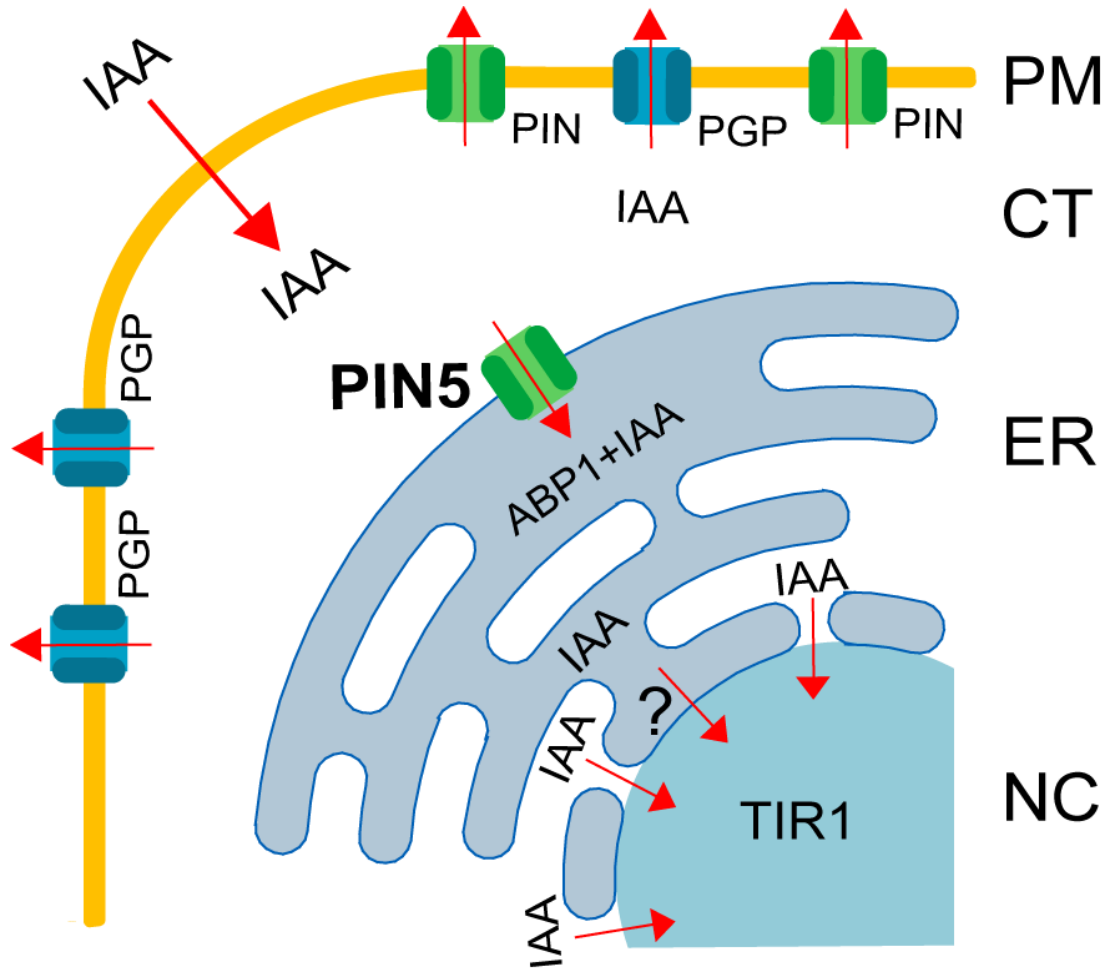




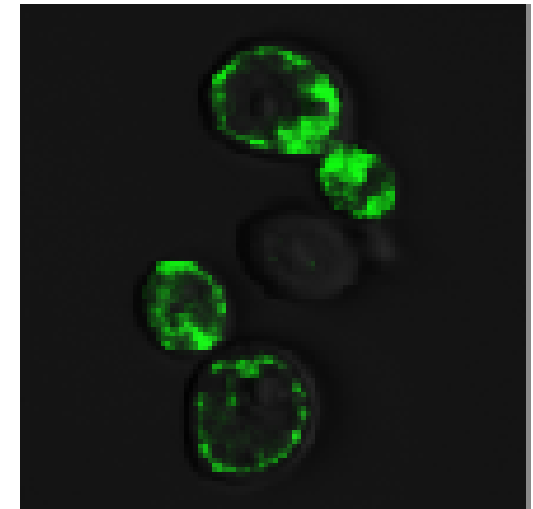
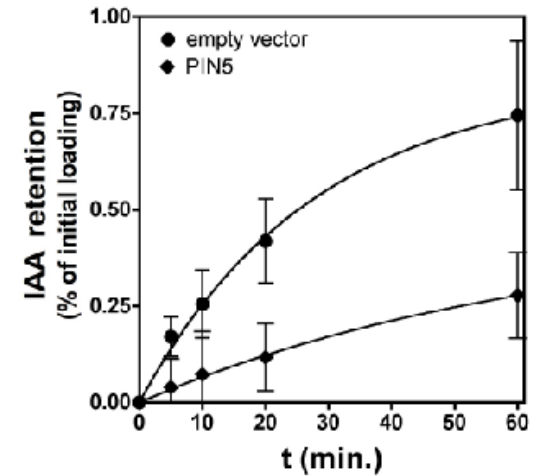
PIN5 regulates auxin metabolism



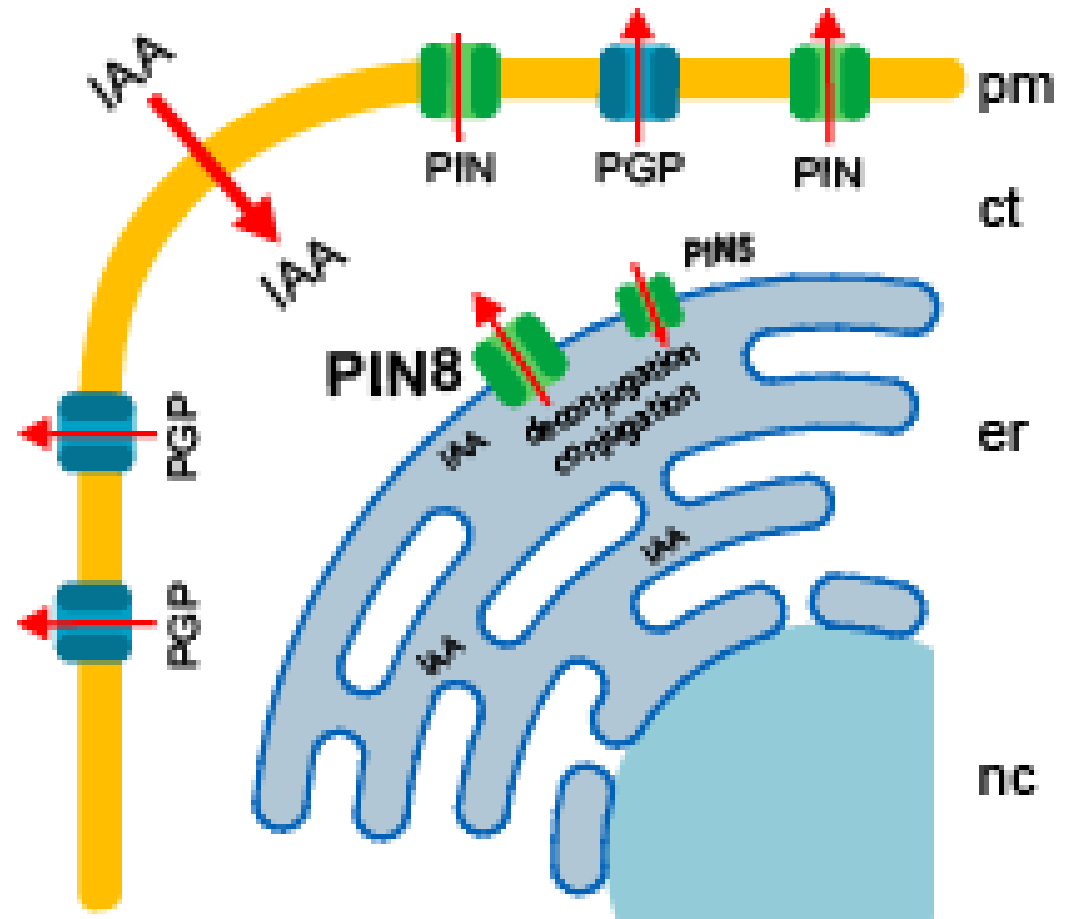
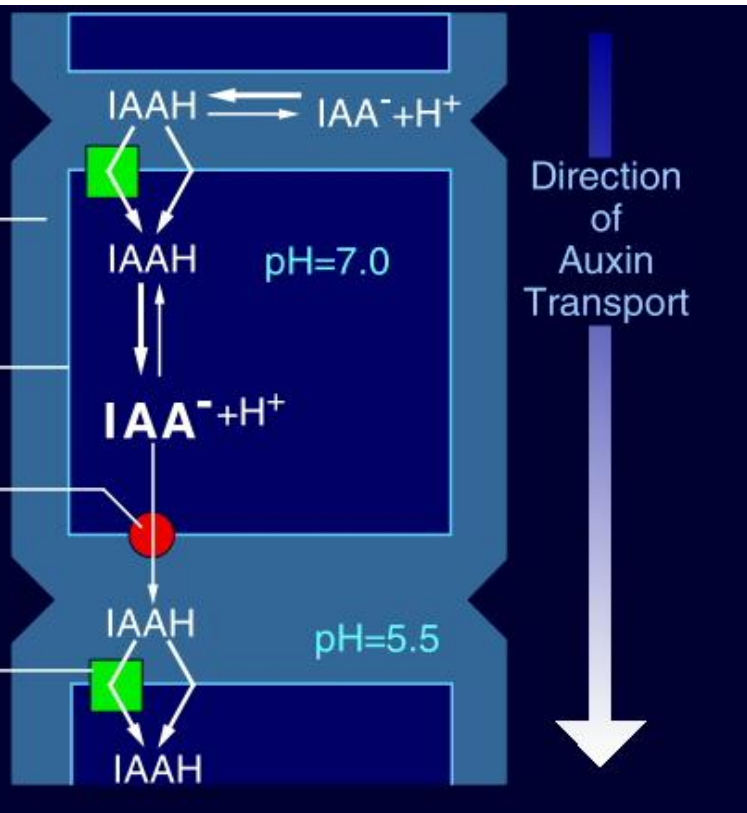
PIN5-dependent auxin transport into ER



Yeast



Updated model for auxin transport



-Interested in novel players in auxin homeostasis

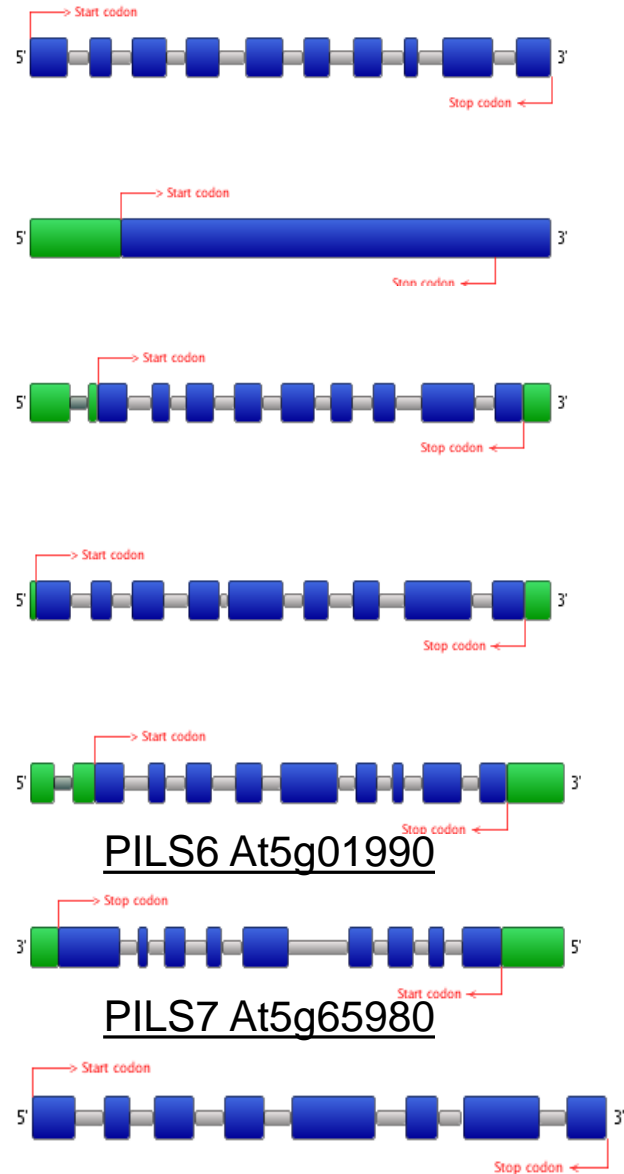
- Search for proteins with PIN like protein organisation

- SMART tool from EMBL

(→finds proteins with topology of interest)

| <u>Atg-code</u> | gene name |
|-----------------|--------------|
| At1g20925 | <i>PILS1</i> |
| At1g71090 | <i>PILS2</i> |
| At1g76520 | <i>PILS3</i> |
| At1g76530 | <i>PILS4</i> |
| At2g17500 | <i>PILS5</i> |
| At5g01990 | <i>PILS6</i> |
| At5g65980 | <i>PILS7</i> |

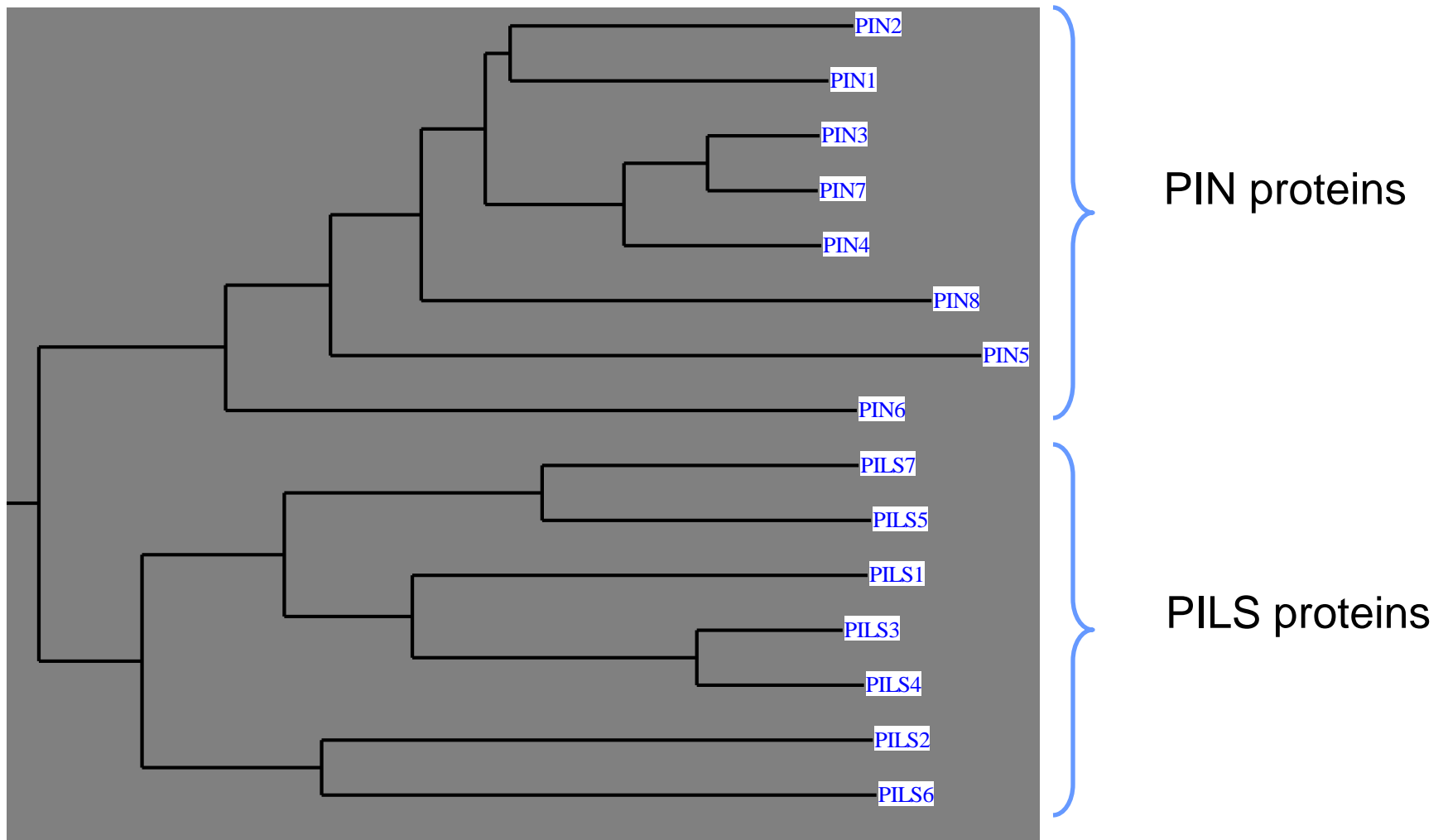
The PILS (PIN-likes) gene family



Aim of the project: Initial characterisation of the PILS family members

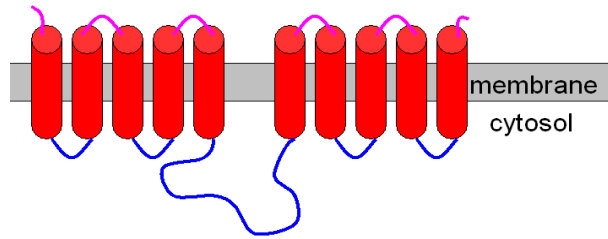
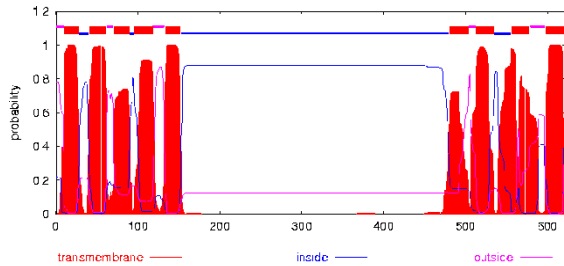
PINs vs PILS: Protein sequences

Limited AA sequence similarities reveal that PINs and PILS form two distinct protein families

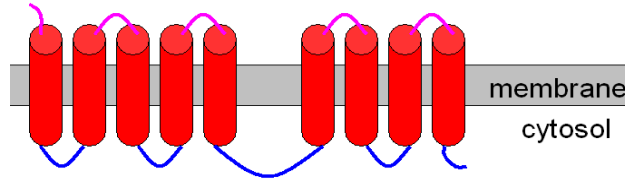
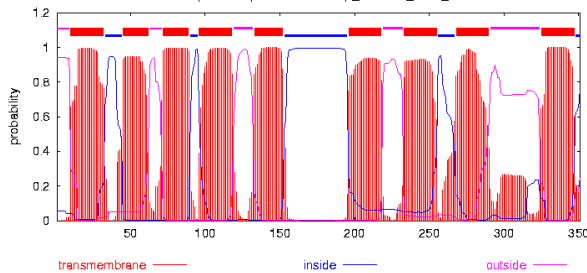


PINs vs PILS: Topology

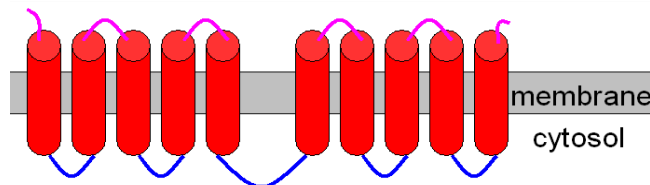
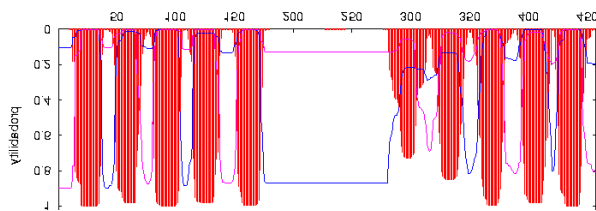
- PILS proteins share the same predicted protein topology as the PINs
- The short central hydrophylic loop make the PILS more related to the Class I PINs



PIN1 (ClassII PINs):
- Long hydrophylic loop



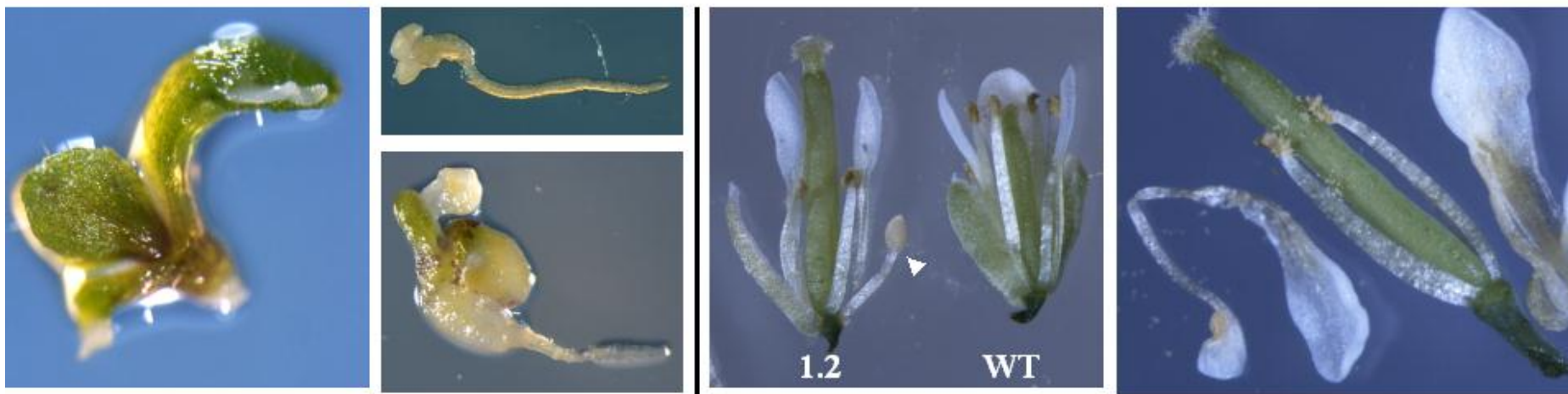
PIN8 (ClassI PINs):
- Short hydrophylic loop



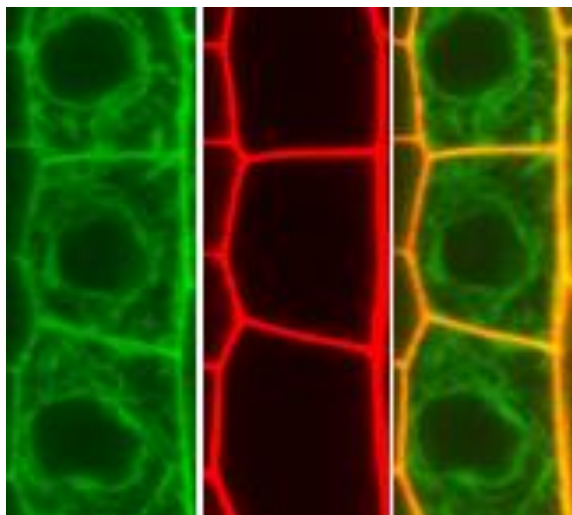
PILS2
- Short hydrophylic loop

PILS genes in Arabidopsis

Phenotypes of *pils* mutants



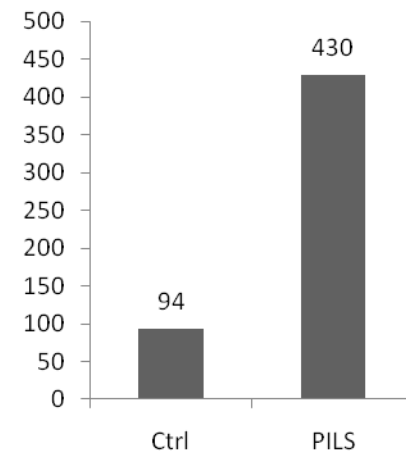
ER Localization



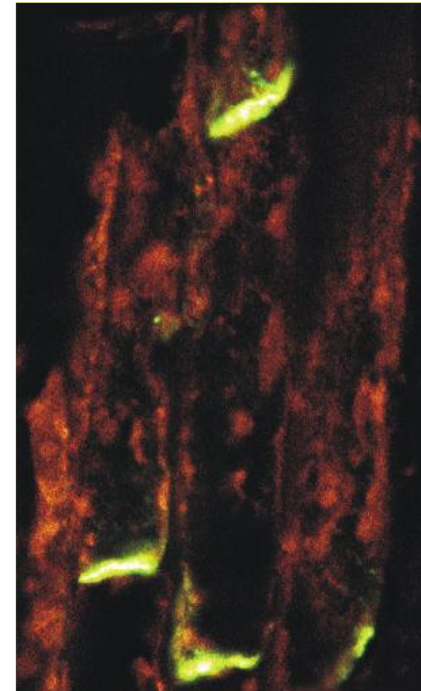
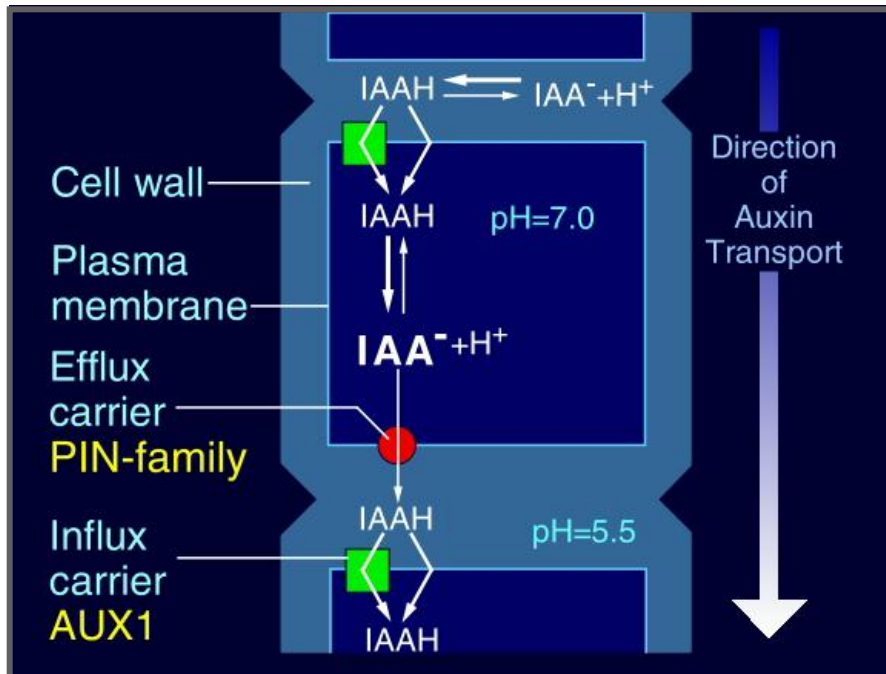
Yeast: IAA transport



IAA Retention

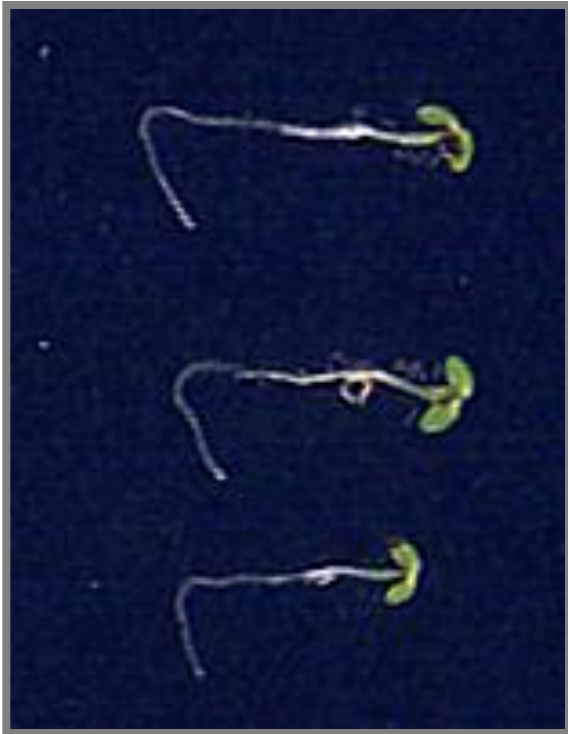


Cellular Polarity of PIN Localization and Directionality of Intercellular Auxin Flow

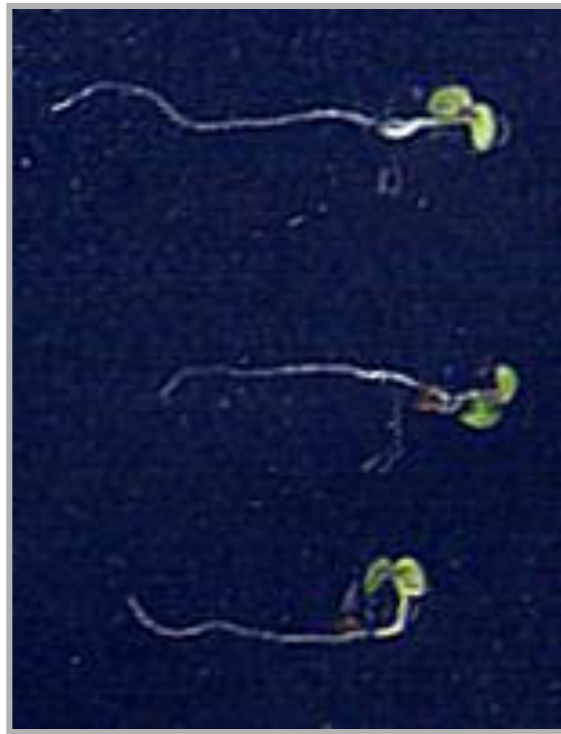


PIN2 – Root Gravitropism

Col-0



pin2



PIN2 protein

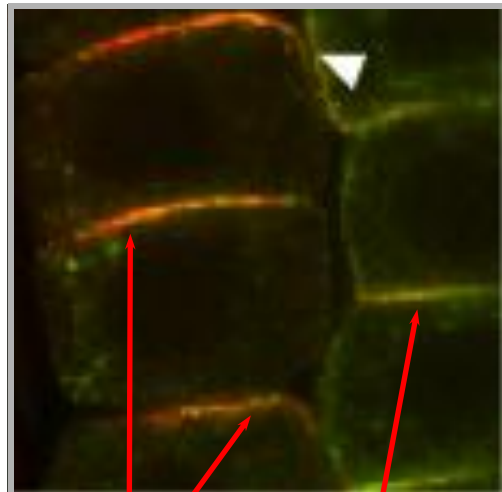


PIN-specific Signals for Polar Targeting

PIN2pr::PIN2:HA

PIN2pr::PIN1:HA

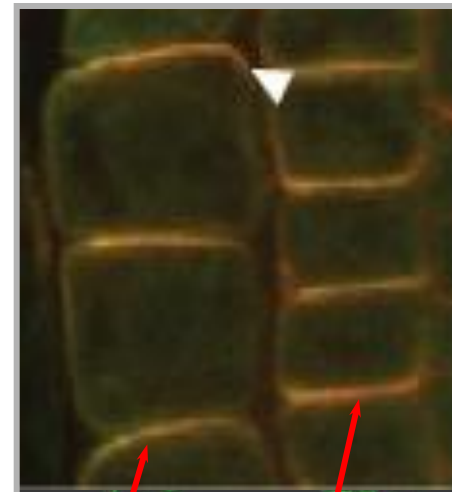
PIN2pr::PIN1:GFP



apical
basal
localization



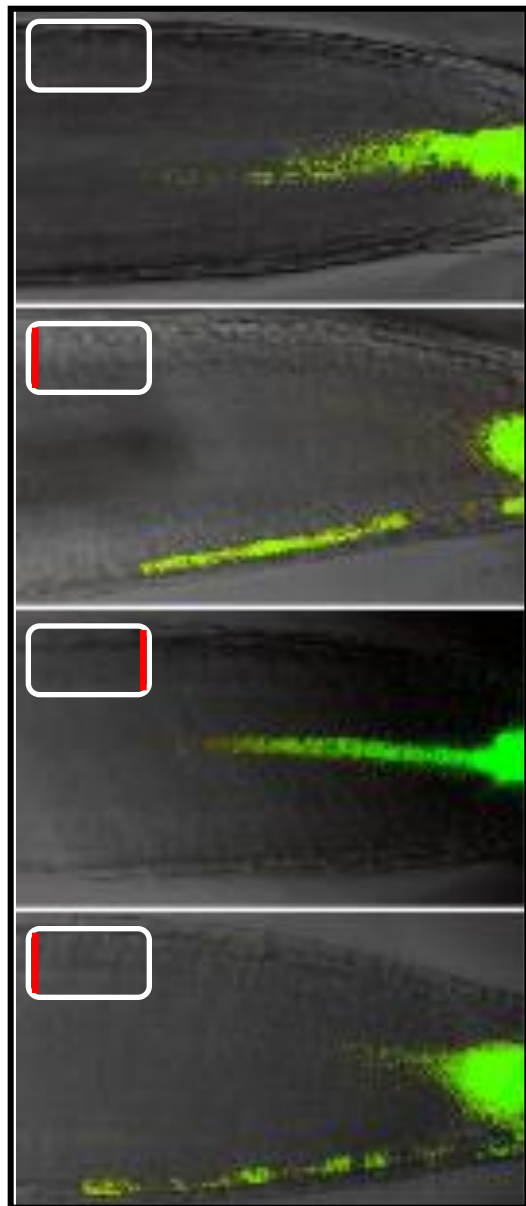
basal
localization



apical
basal
localization

PIN Polarity Determines Direction of Auxin Flow

DR5rev::GFP



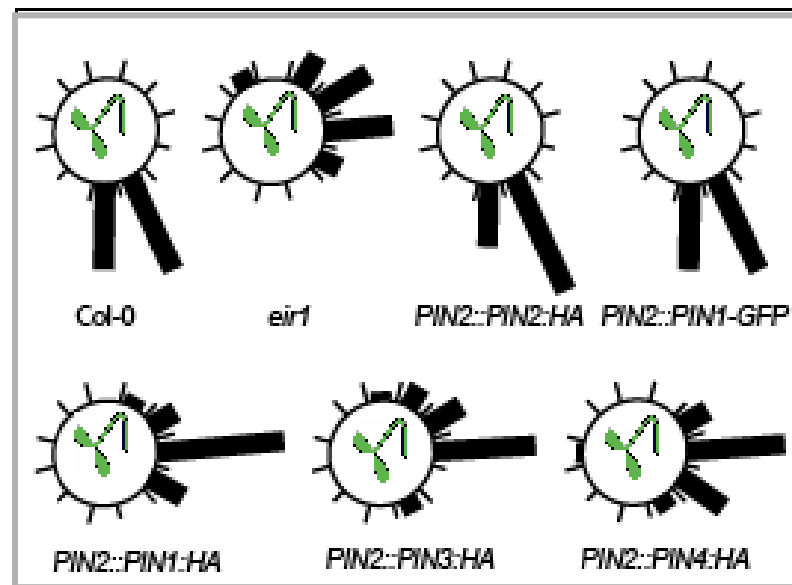
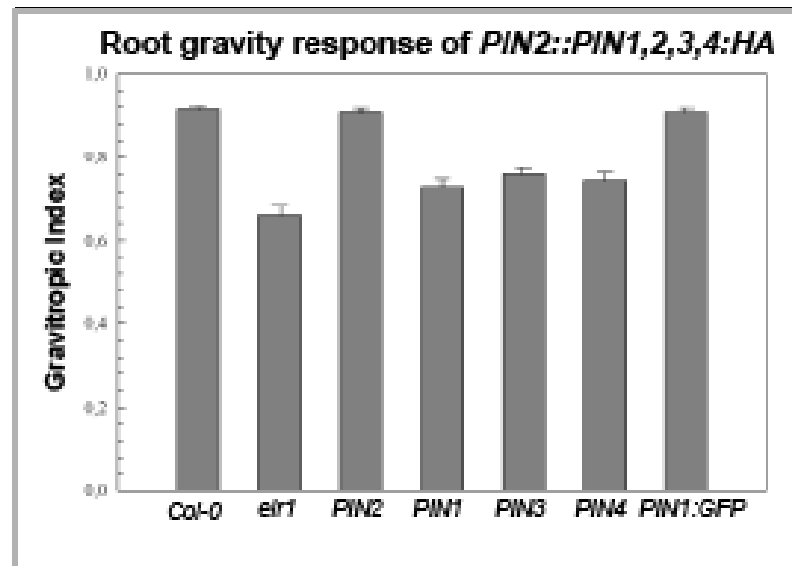
pin2 (eir1, agr1)

PIN2::PIN2:HA

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

PIN2::PIN1:GFP-3

gravitropism



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

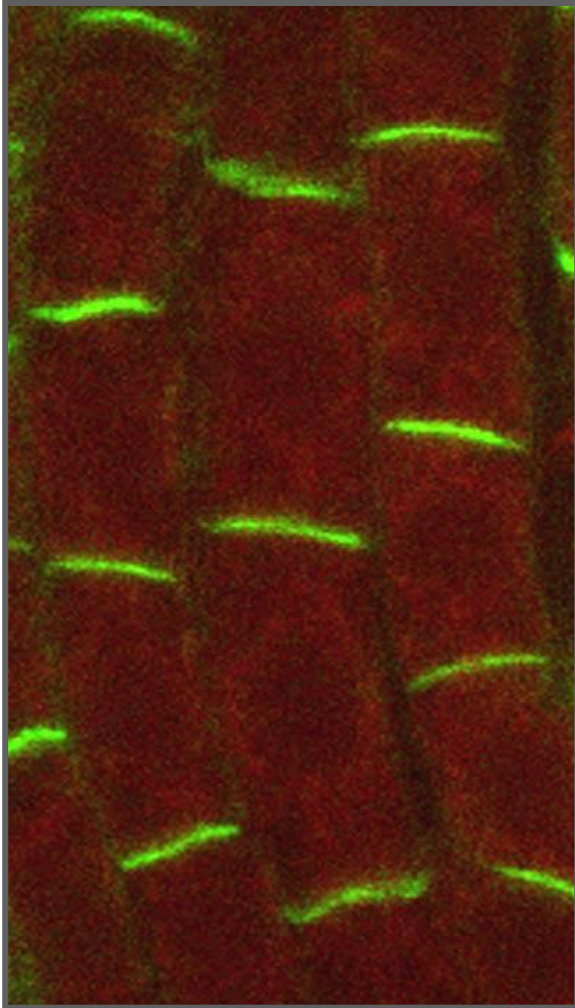
and

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

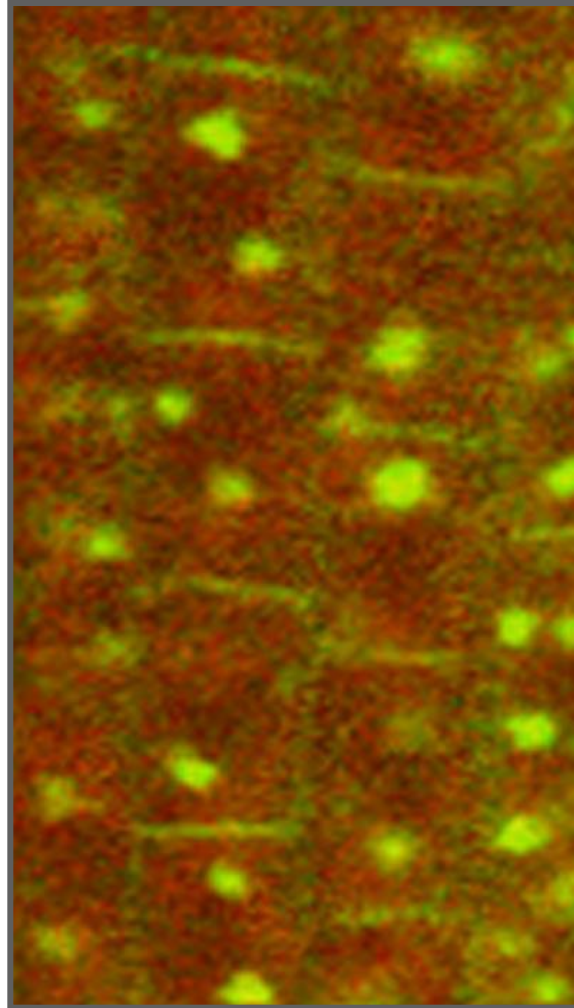
Constitutive Cycling of PINs

PIN1 Subcellular Movement

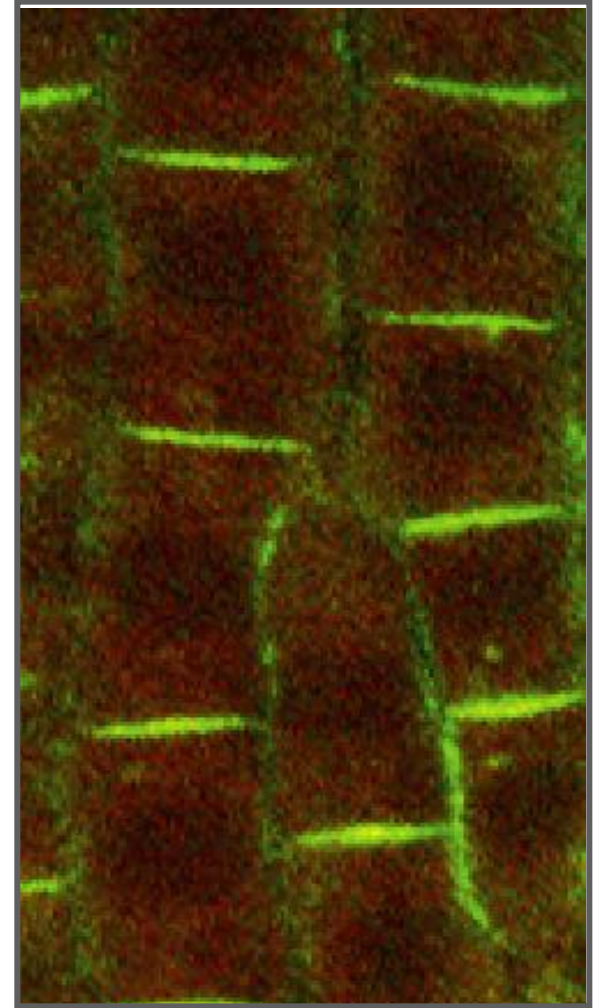
untreated



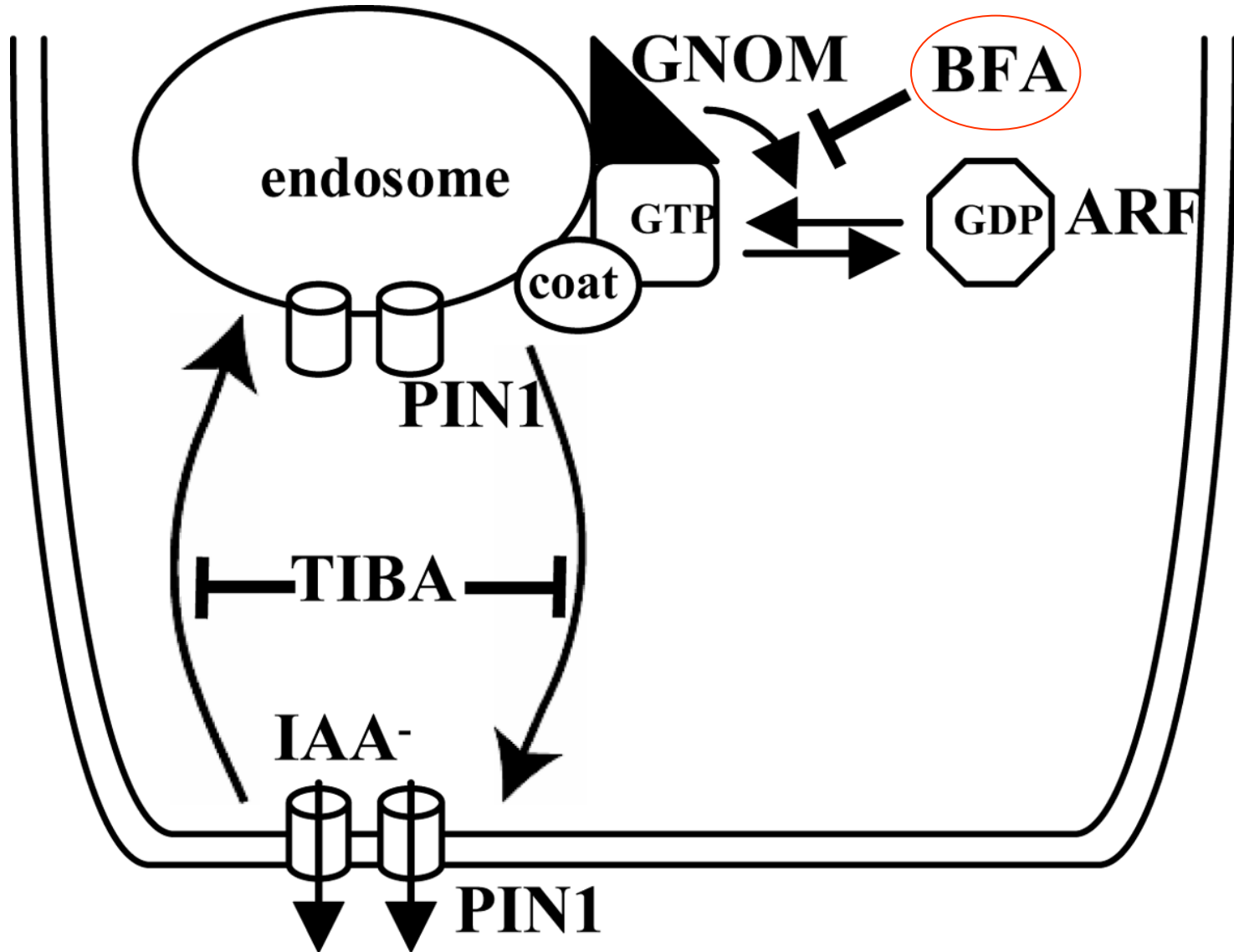
+ *BFA*



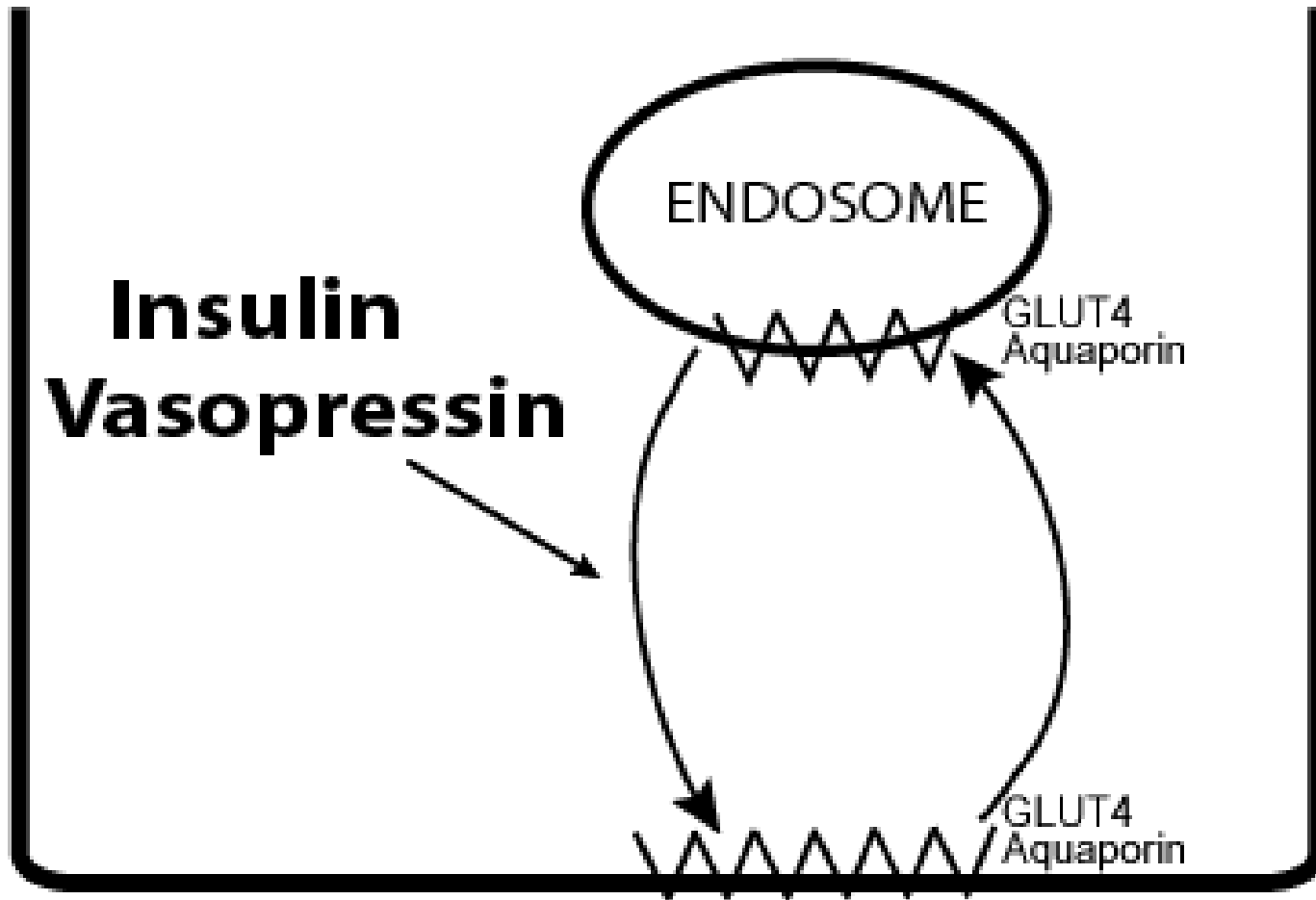
- *BFA*



Dynamic Movement of PIN Proteins

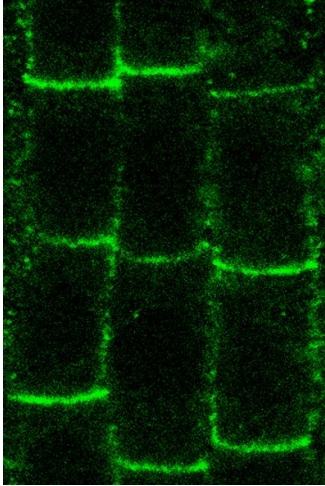


Subcellular Cycling – Means to Modulate Protein Activity?

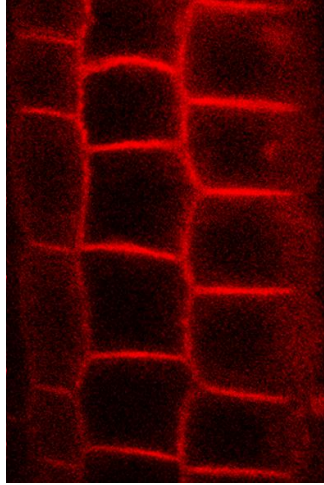


Auxin Inhibits Internalization of Plasma Membrane Proteins

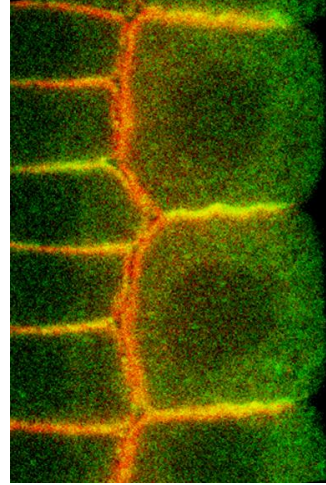
NAA
BFA
/BFA



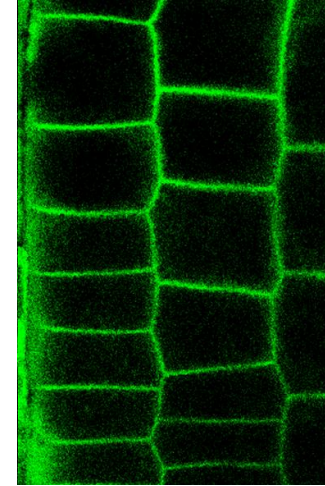
PIN1



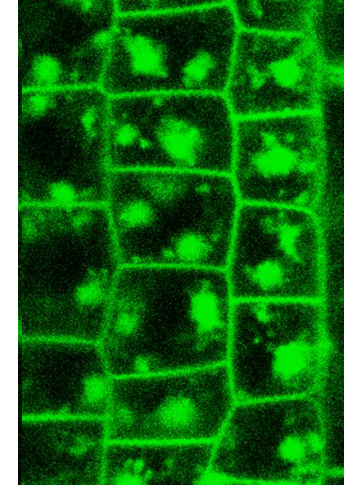
PM-ATPase



PIN2/ATPase

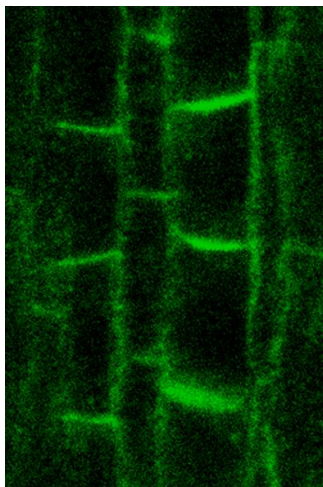


PIP2

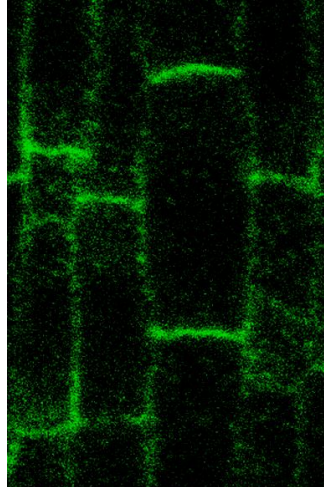


BRI1

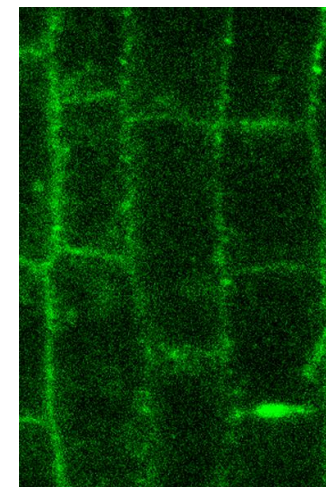
IAA/BFA



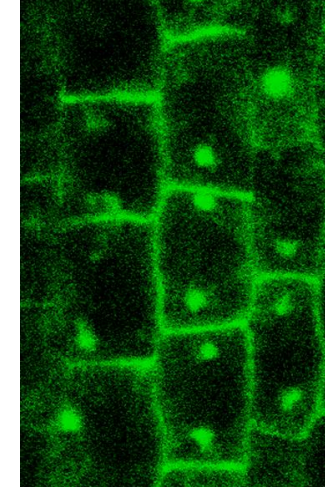
2,4-D/BFA



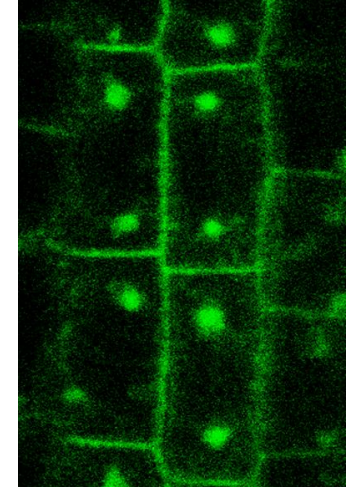
BFA in *sur2*



2-NAA/BFA



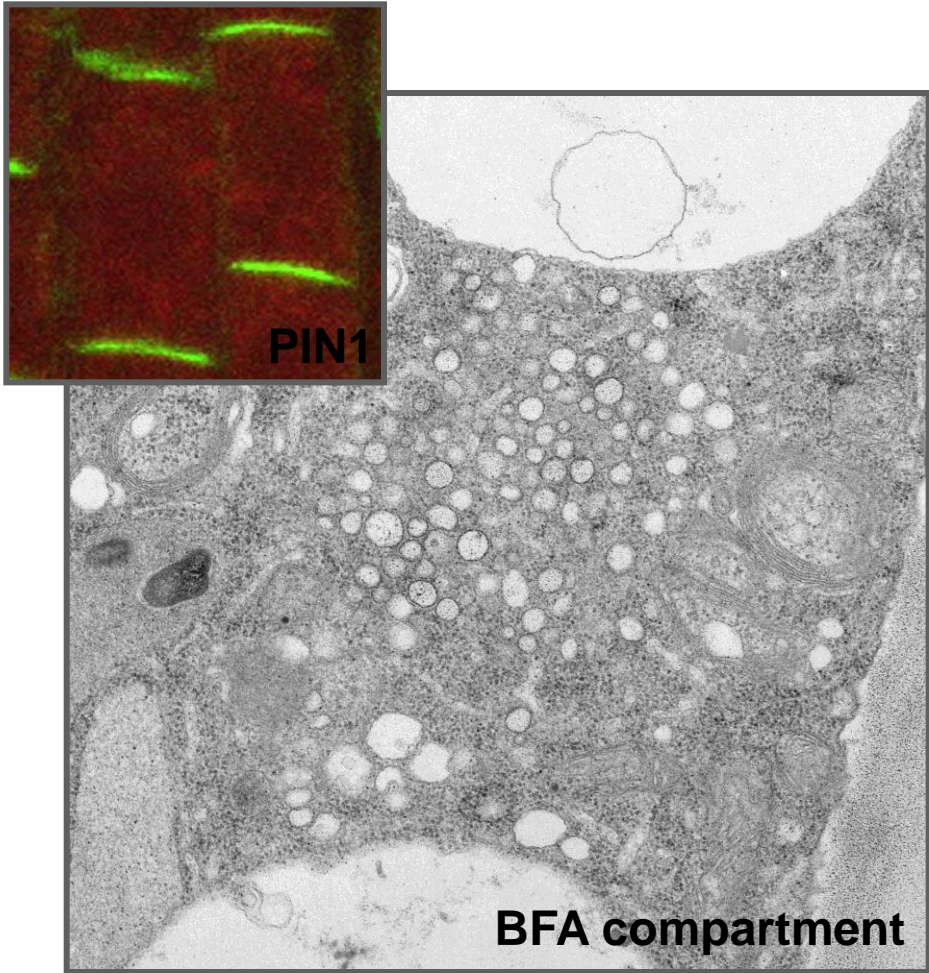
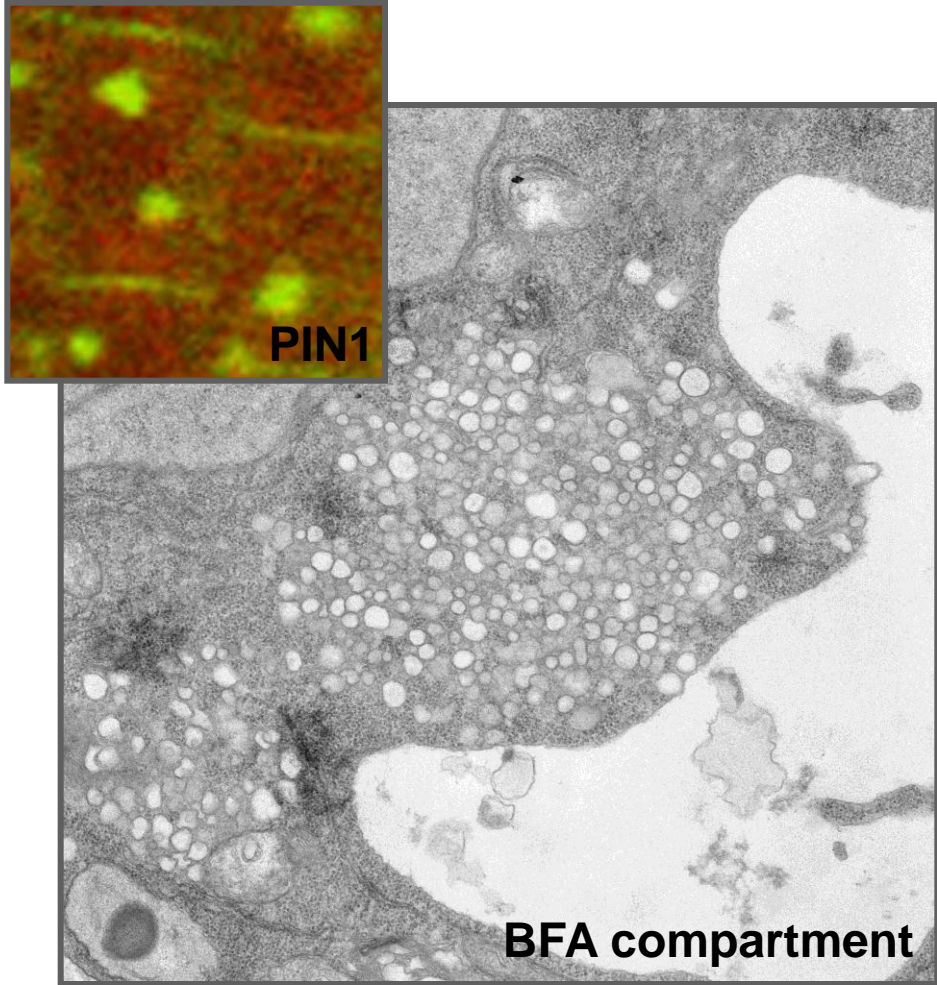
Ethylene/BFA



Place of Auxin Action in Protein Cycling

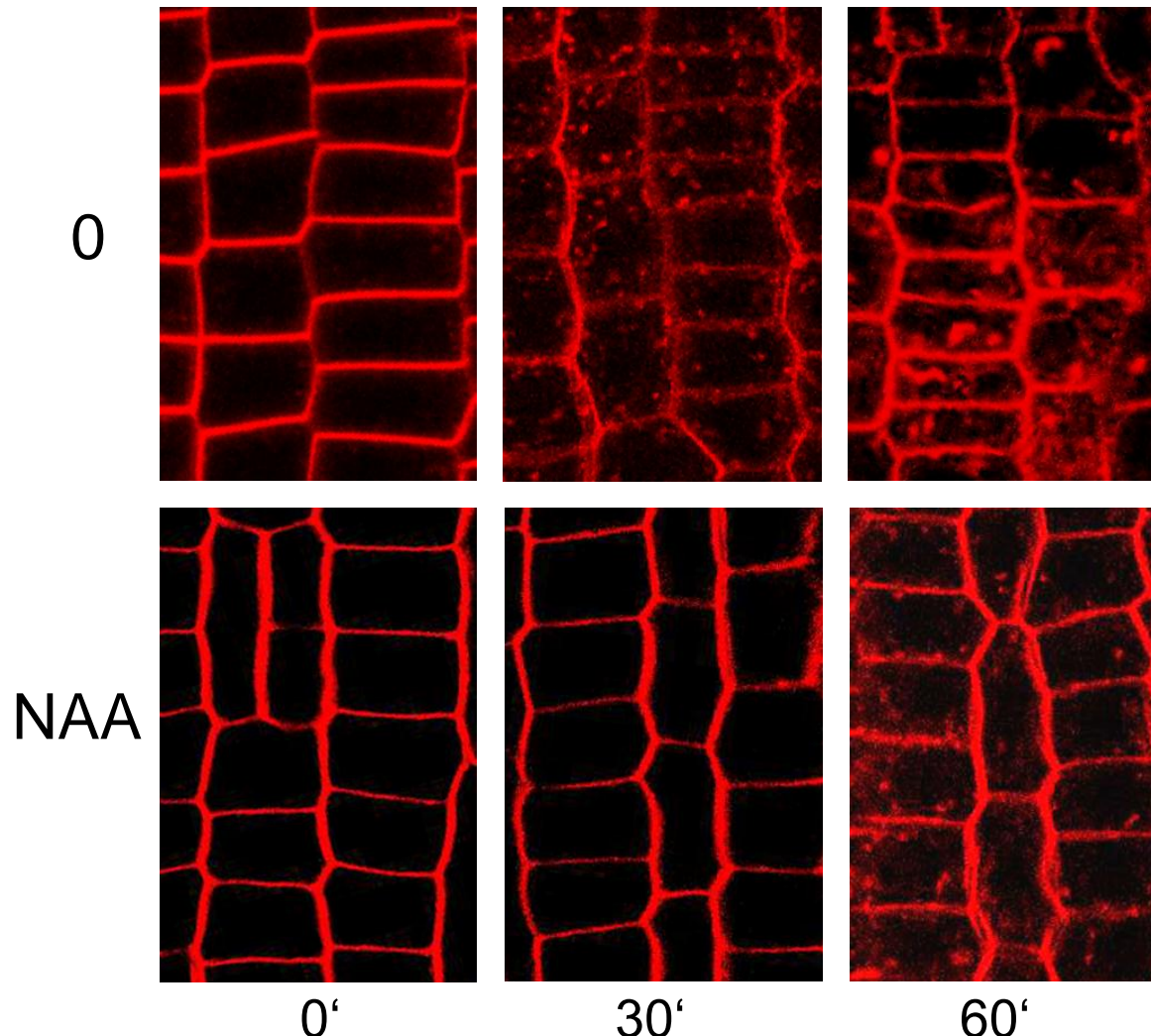
BFA

Auxin + BFA



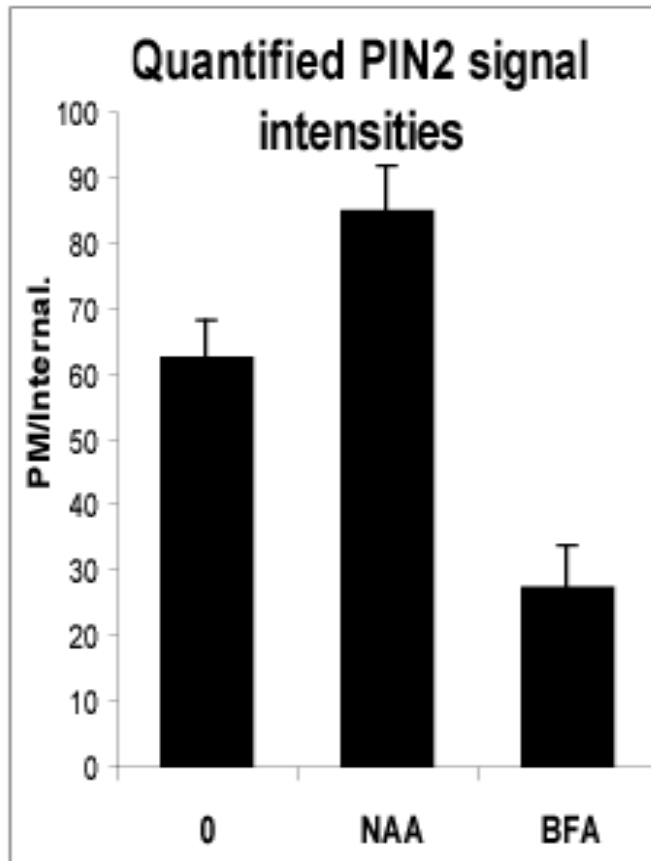
Auxin Inhibits Endocytosis

Uptake of endocytic tracer FM4-64

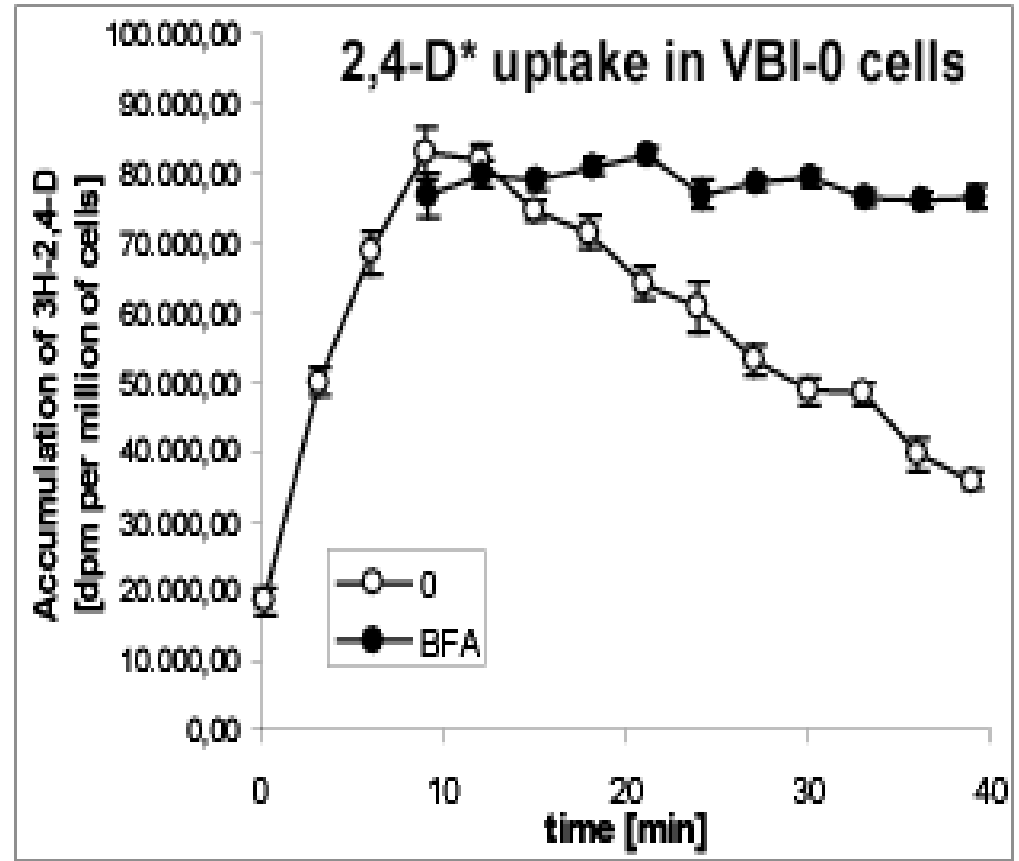


Auxin Increases PIN Levels at Cell Surface and Stimulates its own Efflux

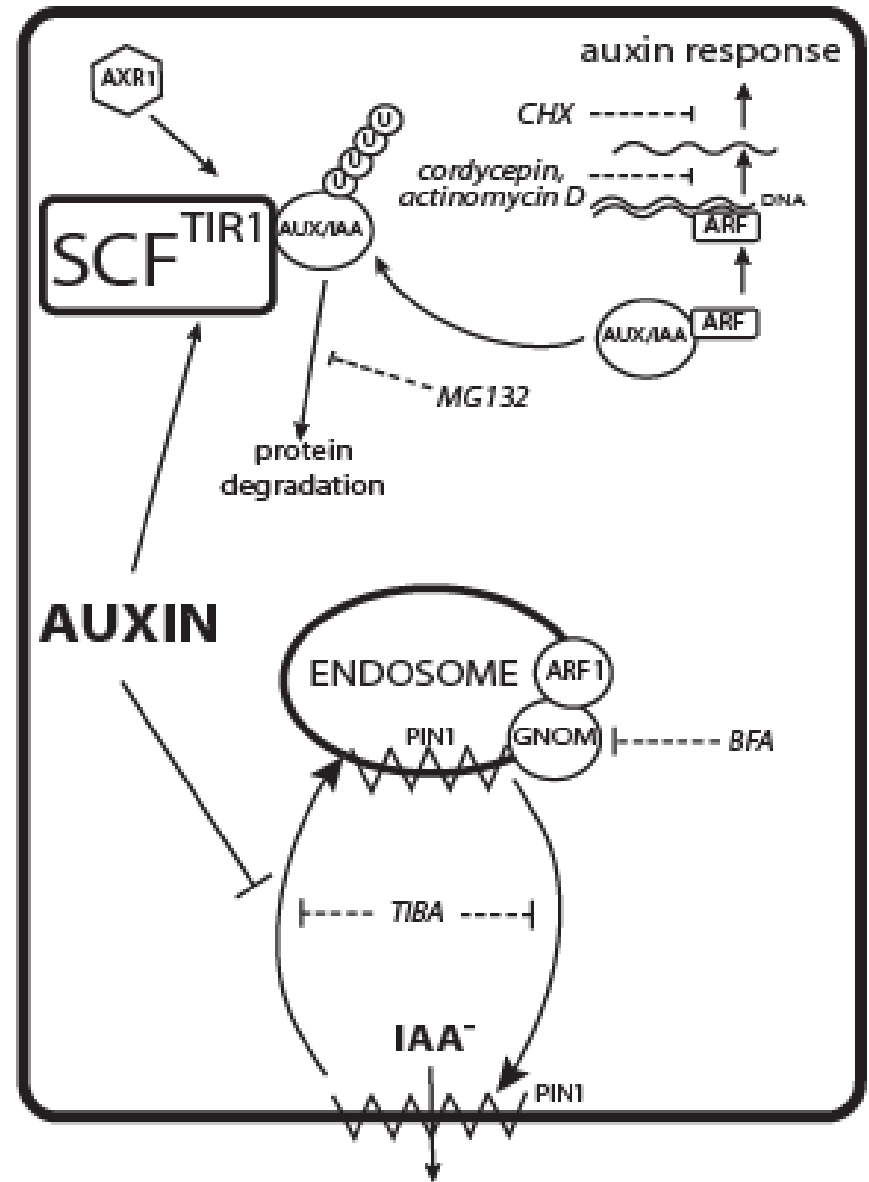
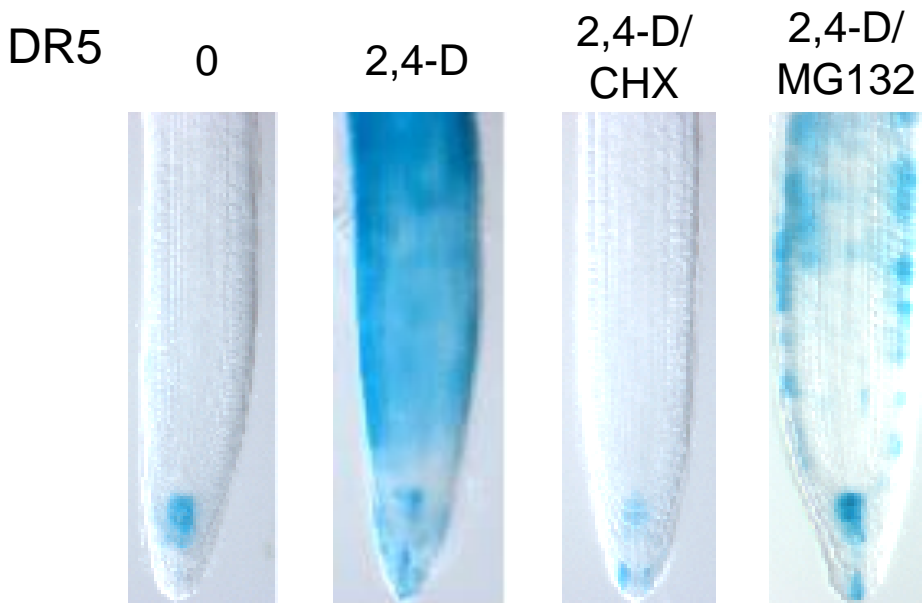
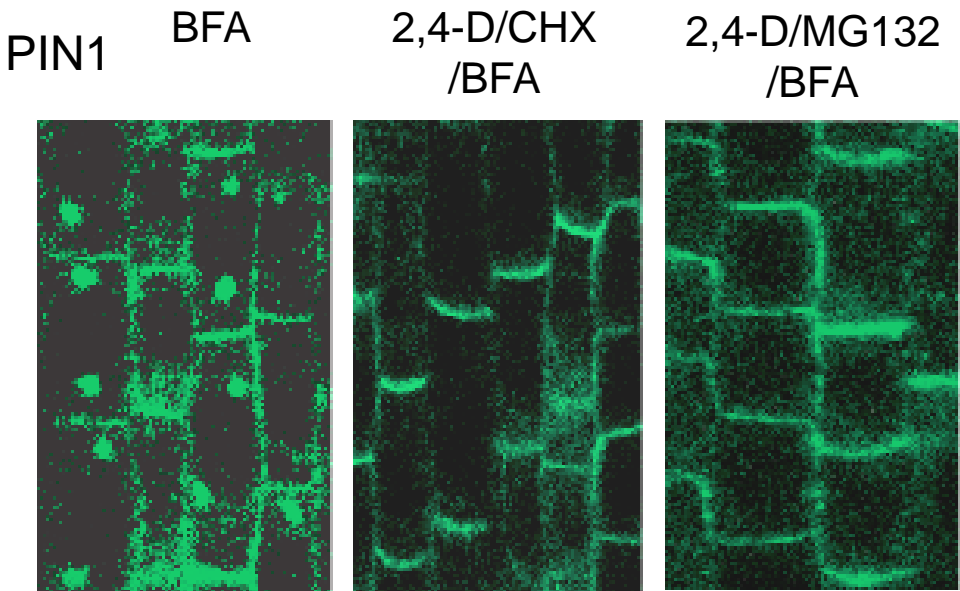
PIN2 levels at PM



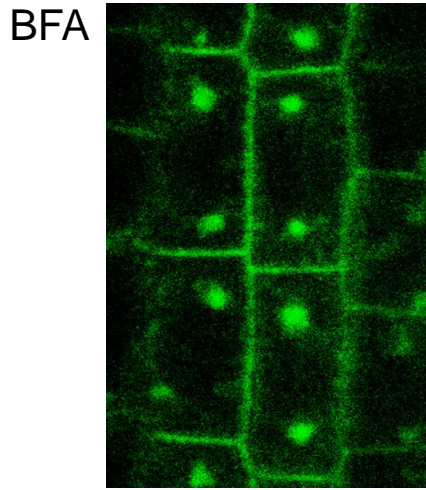
Auxin efflux in tobacco cells



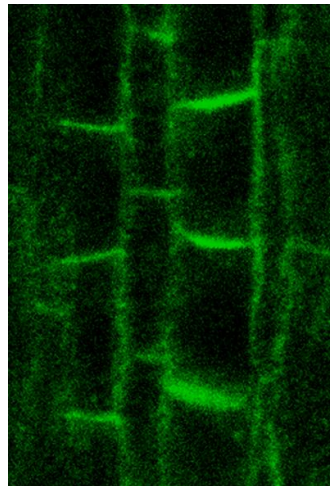
Novel Pathway of Auxin Action



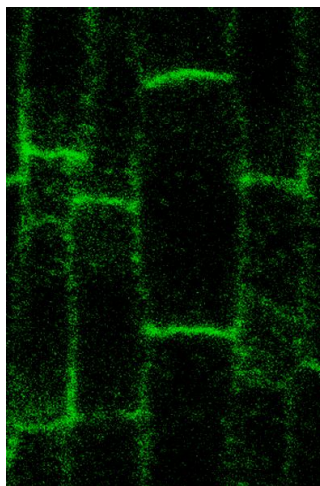
Auxin Inhibits PIN Internalization and Stimulates its Efflux



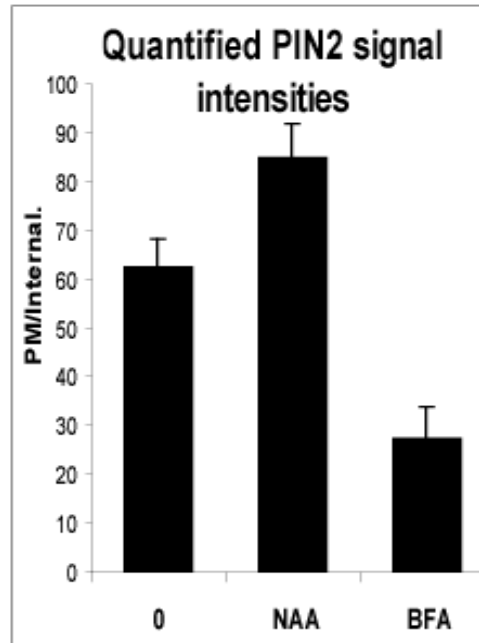
IAA/BFA



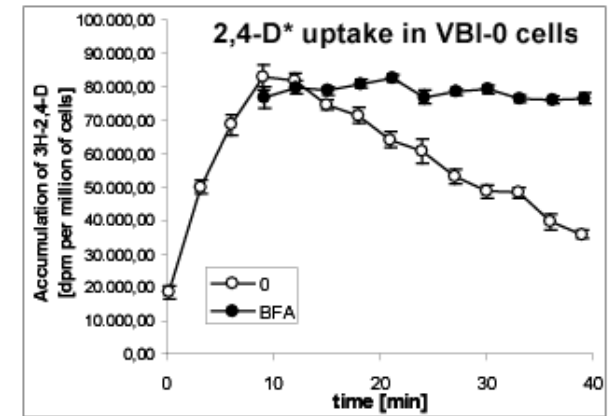
2,4-D/BFA



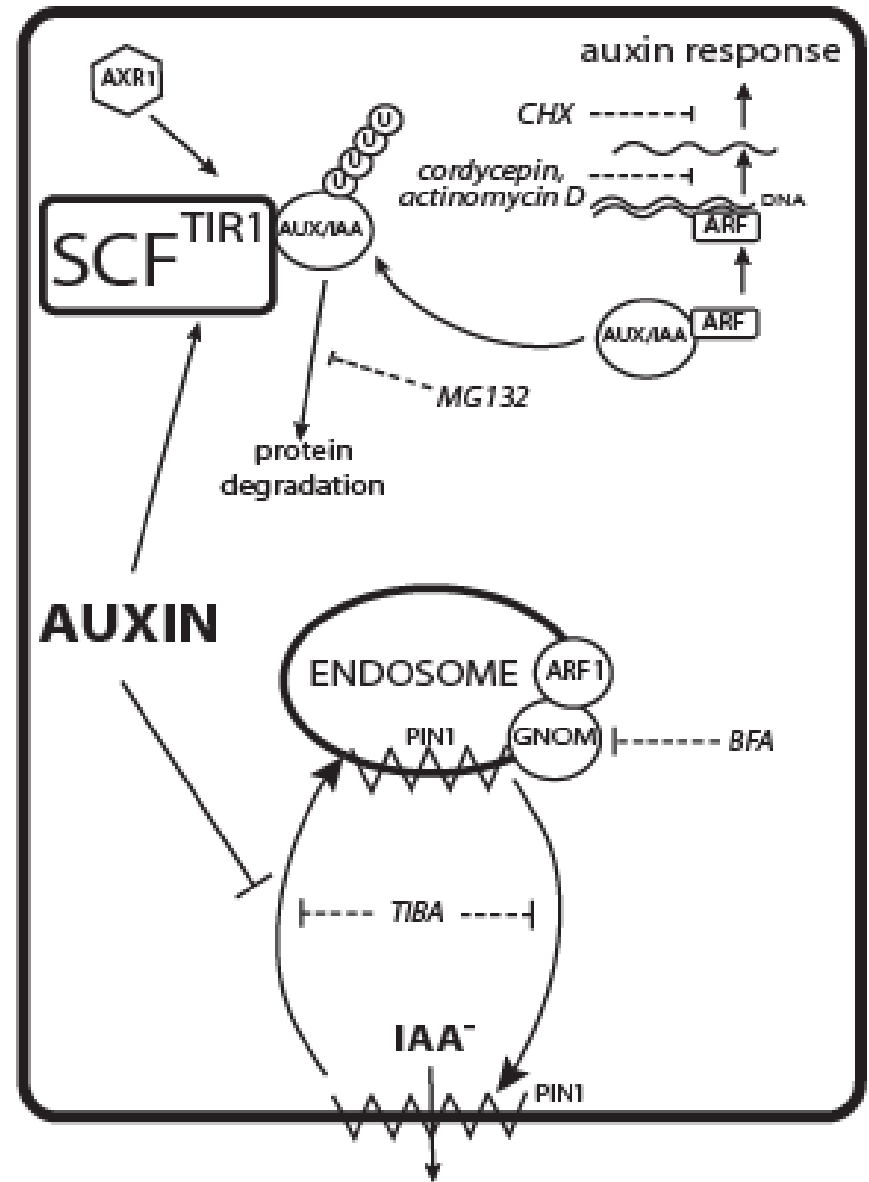
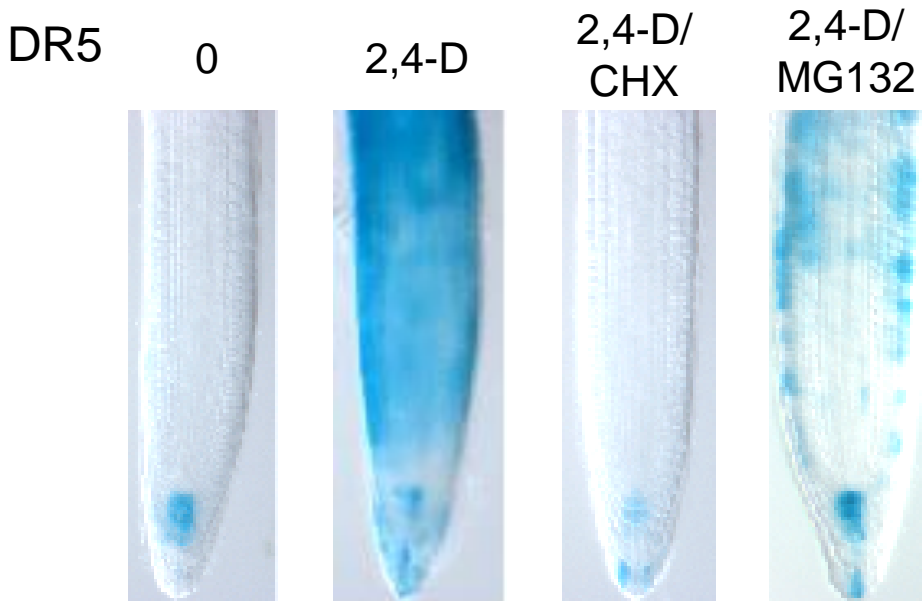
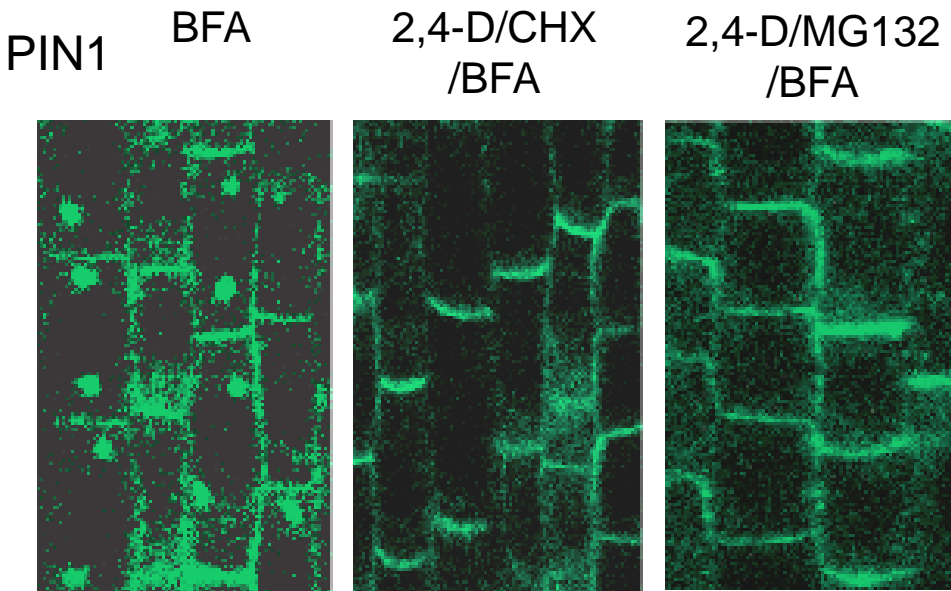
PIN2 at PM



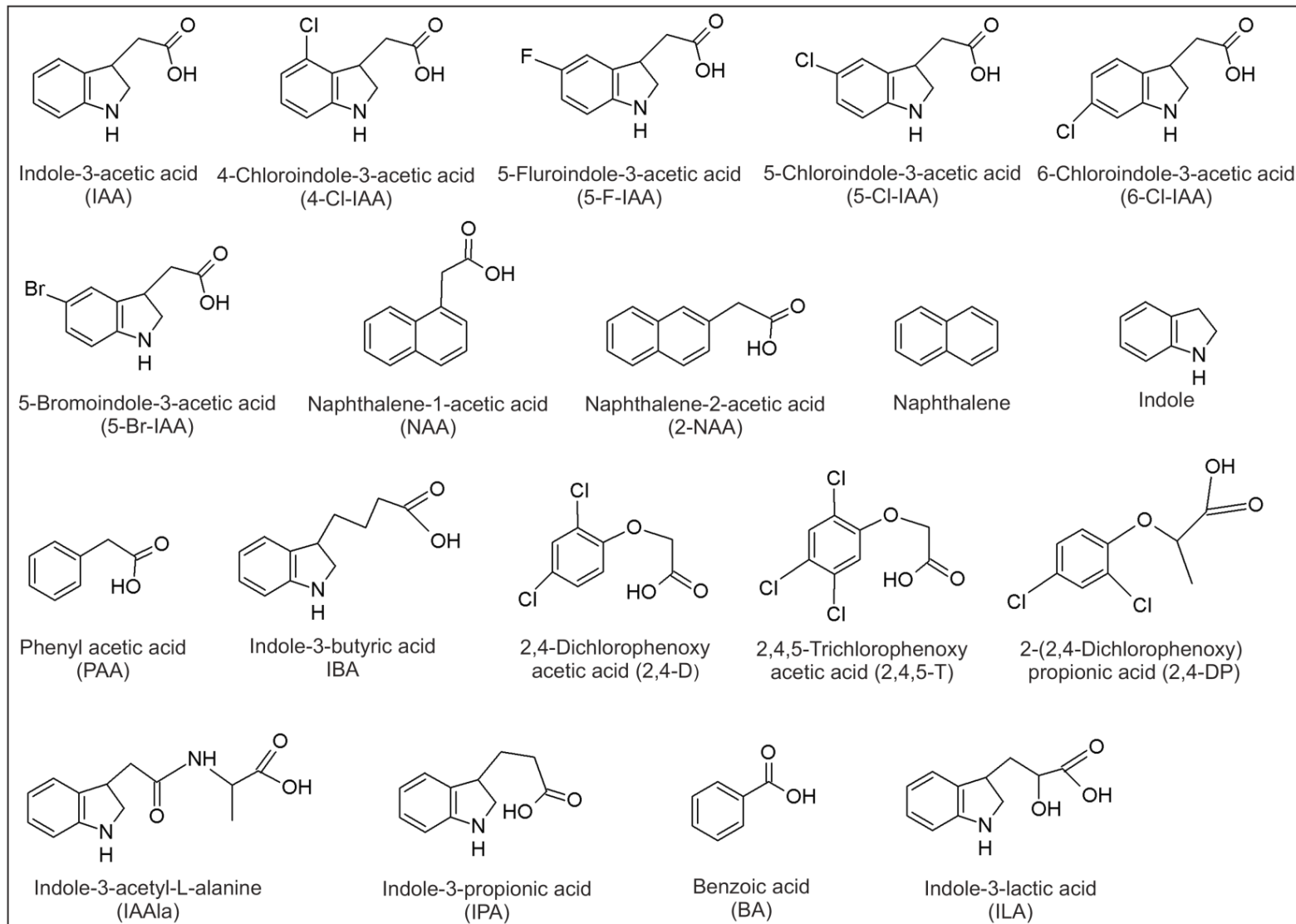
Auxin efflux



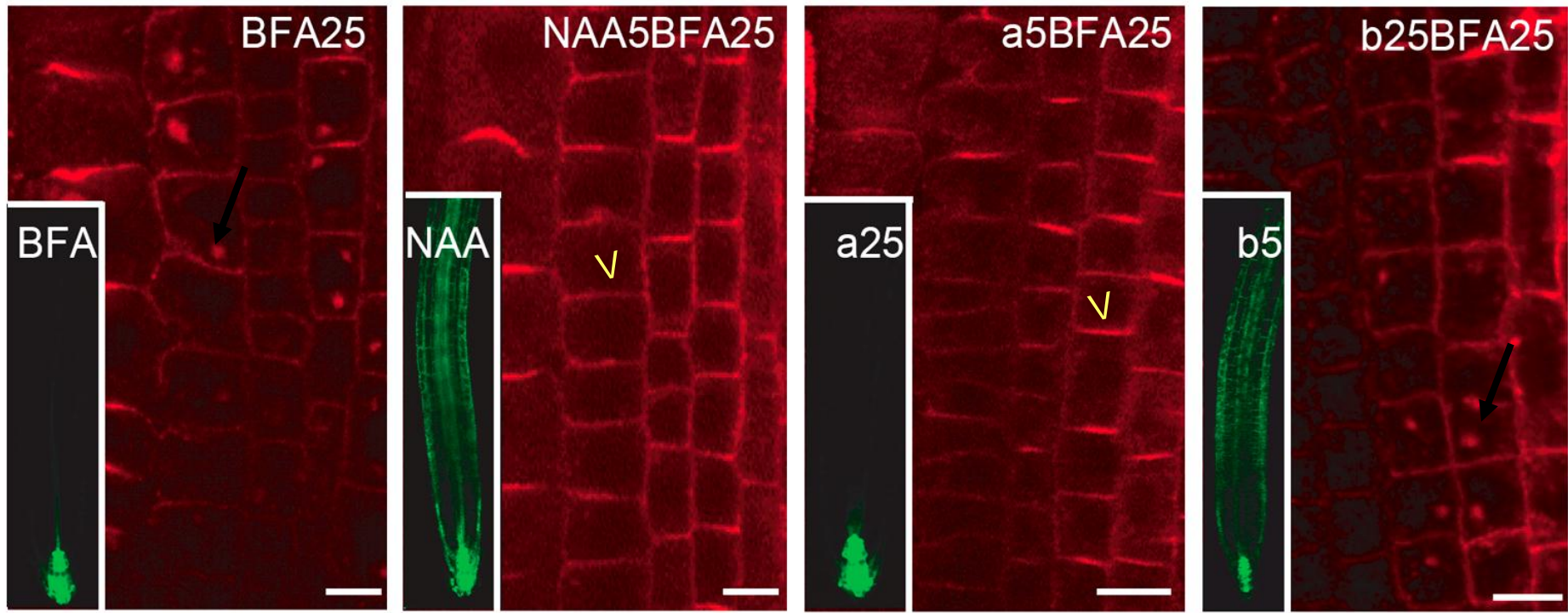
Novel Pathway of Auxin Action



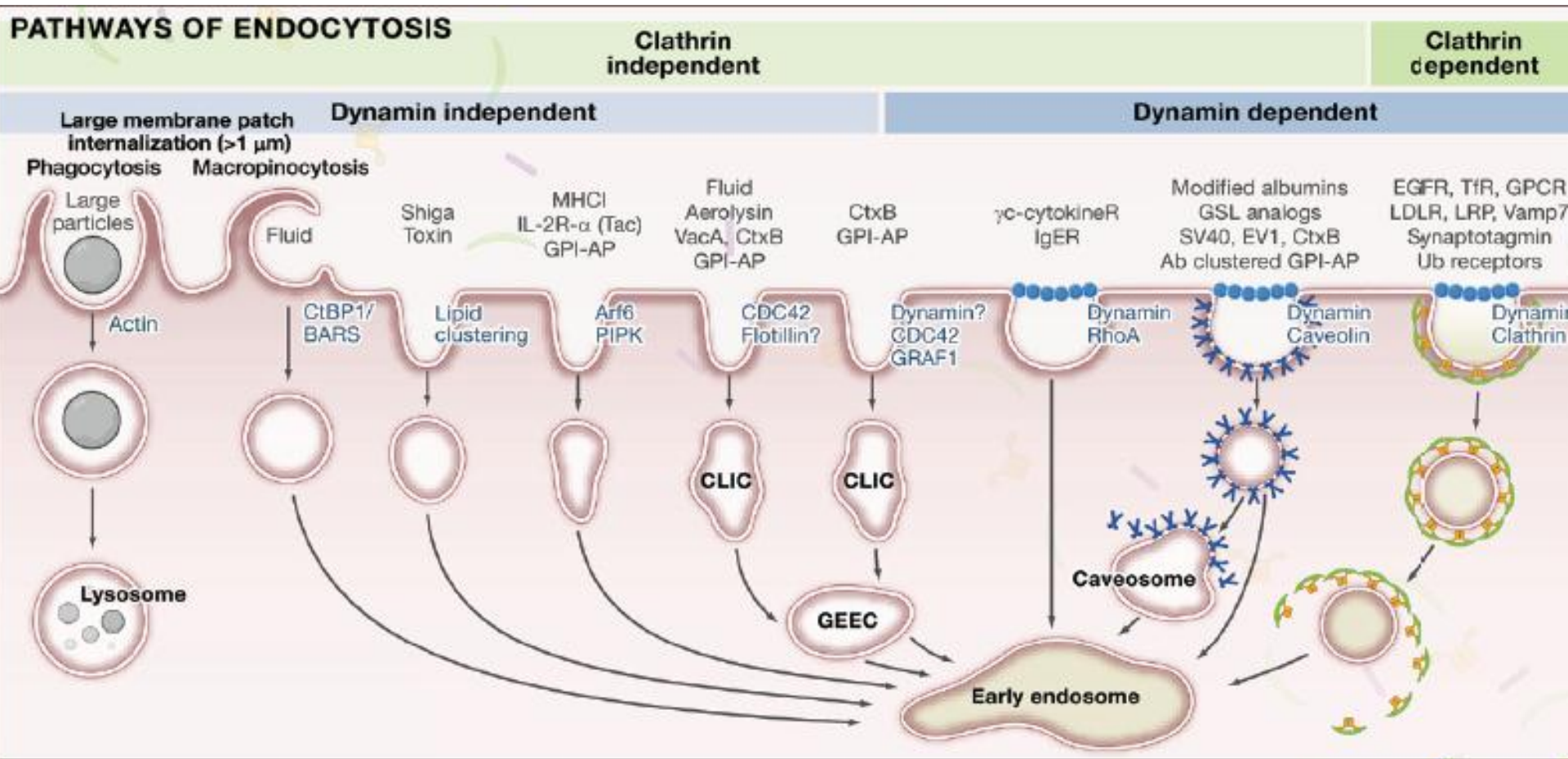
Auxin analogues: mapping the binding sites

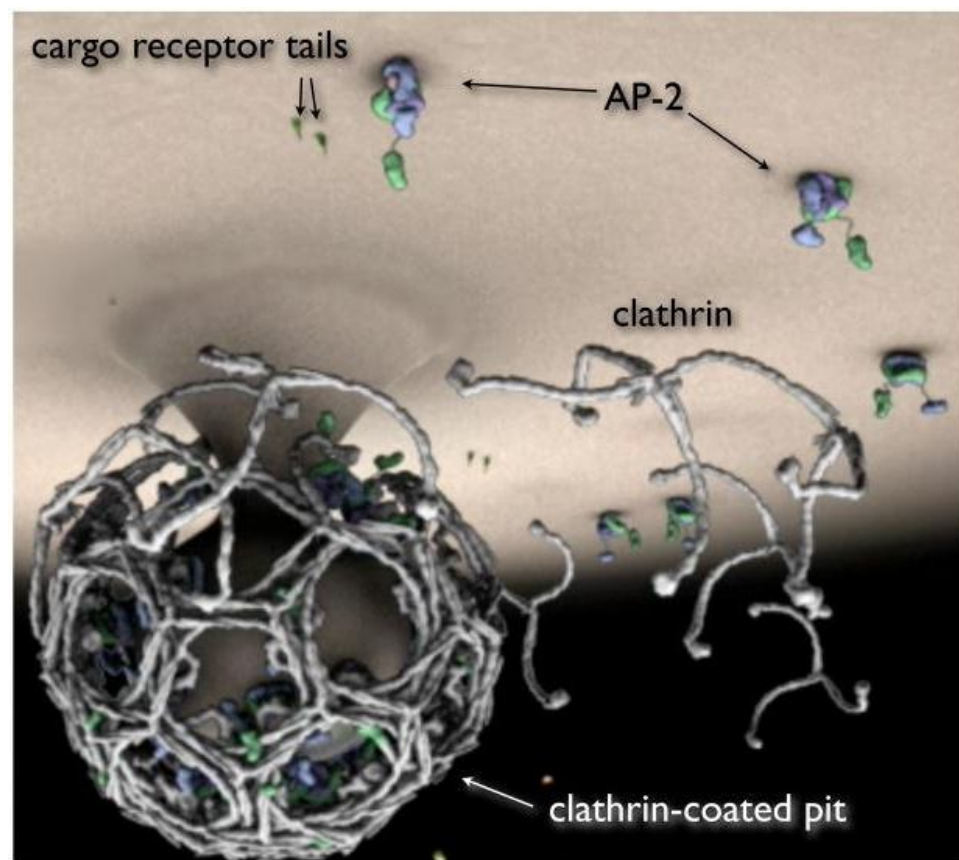
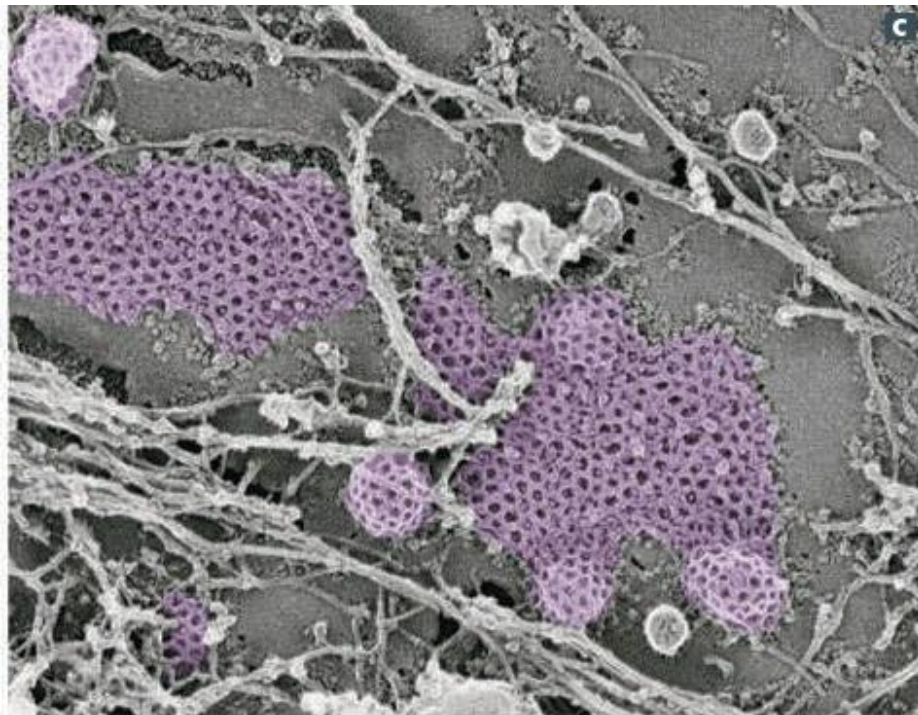


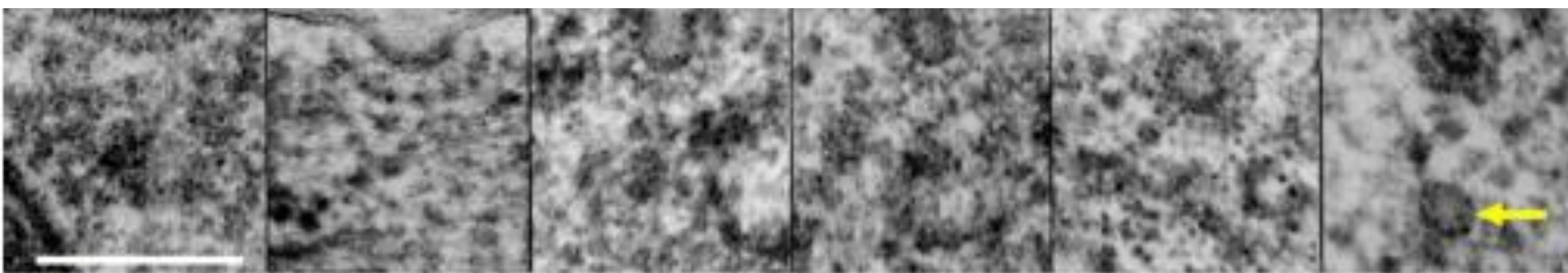
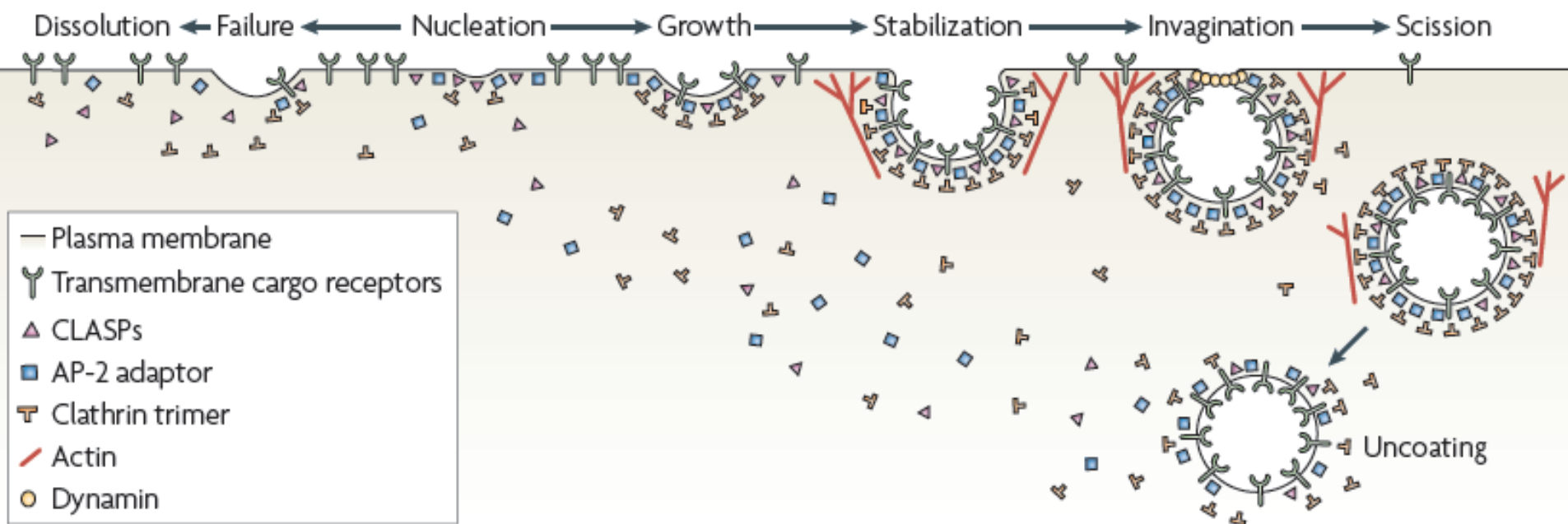
Auxin-mediated regulation of transcription and endocytosis involve different binding sites



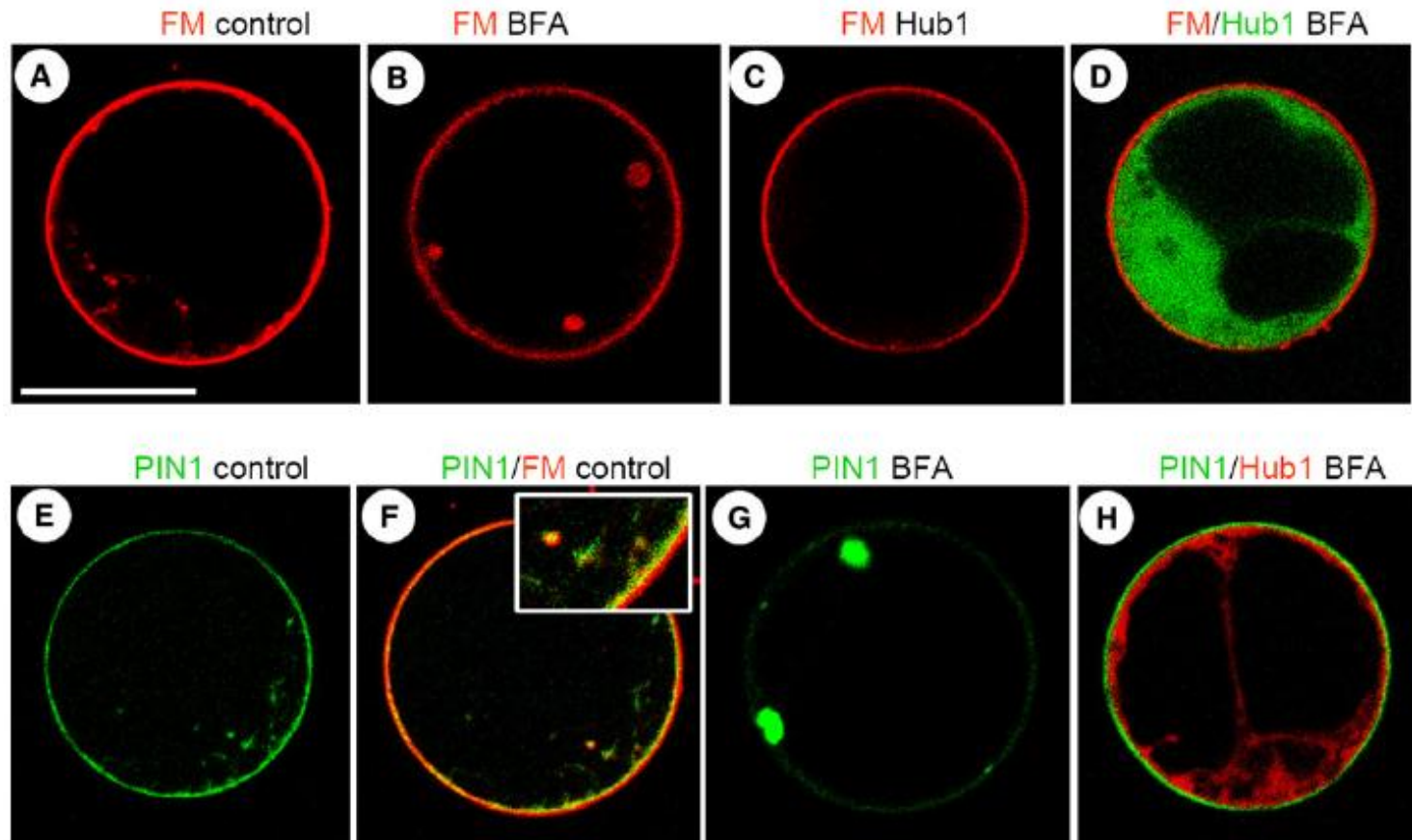
Mechanisms for endocytosis in animals



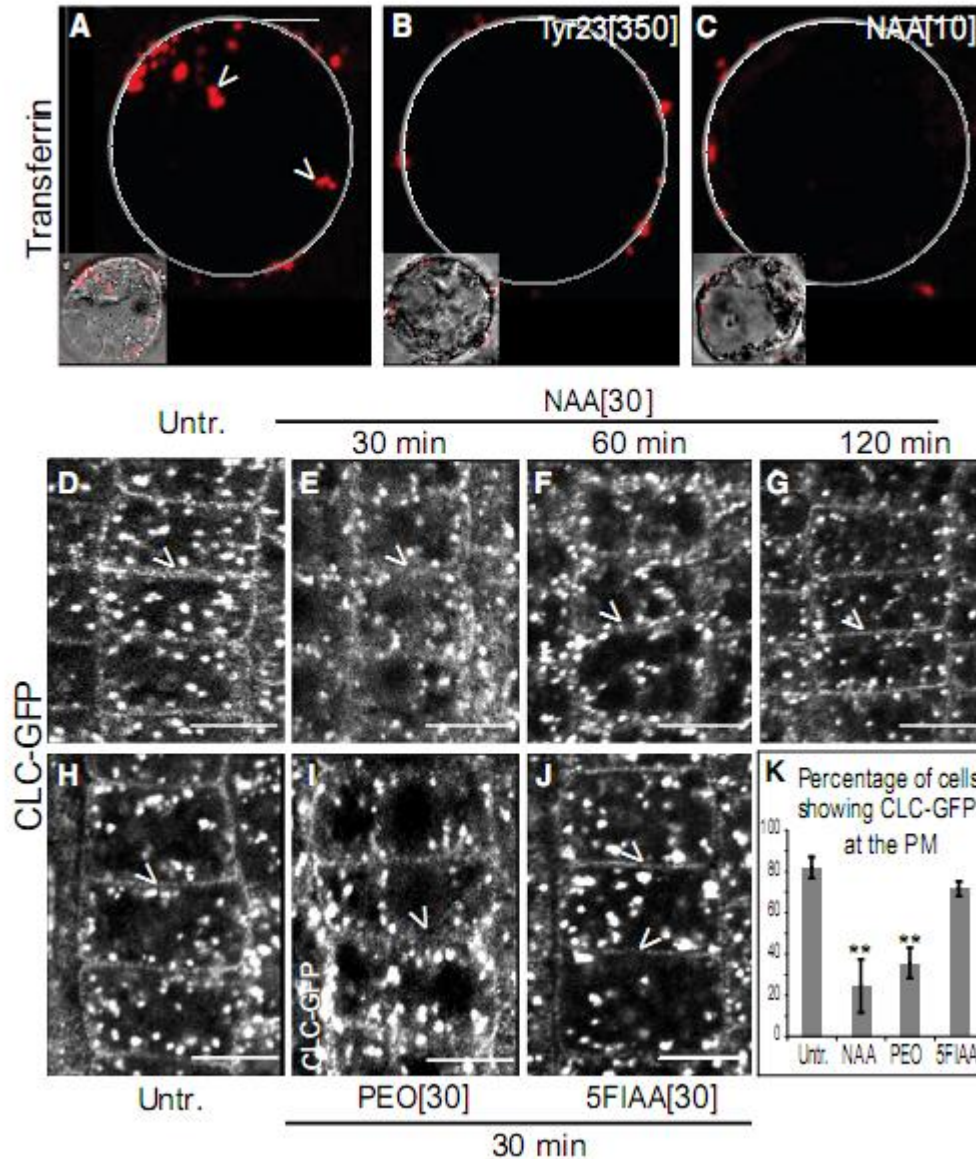




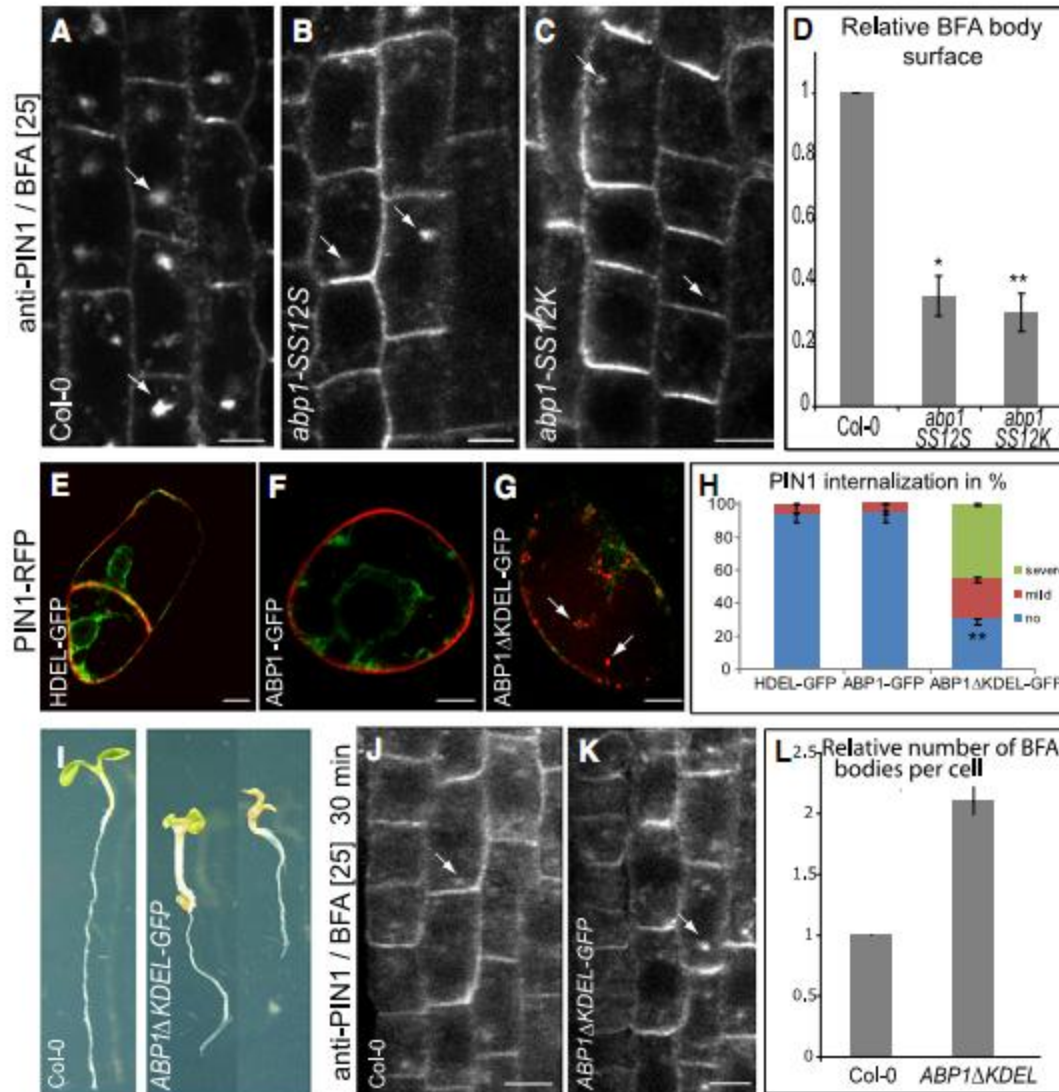
Clathrin is required for PIN internalization in *Arabidopsis* protoplasts



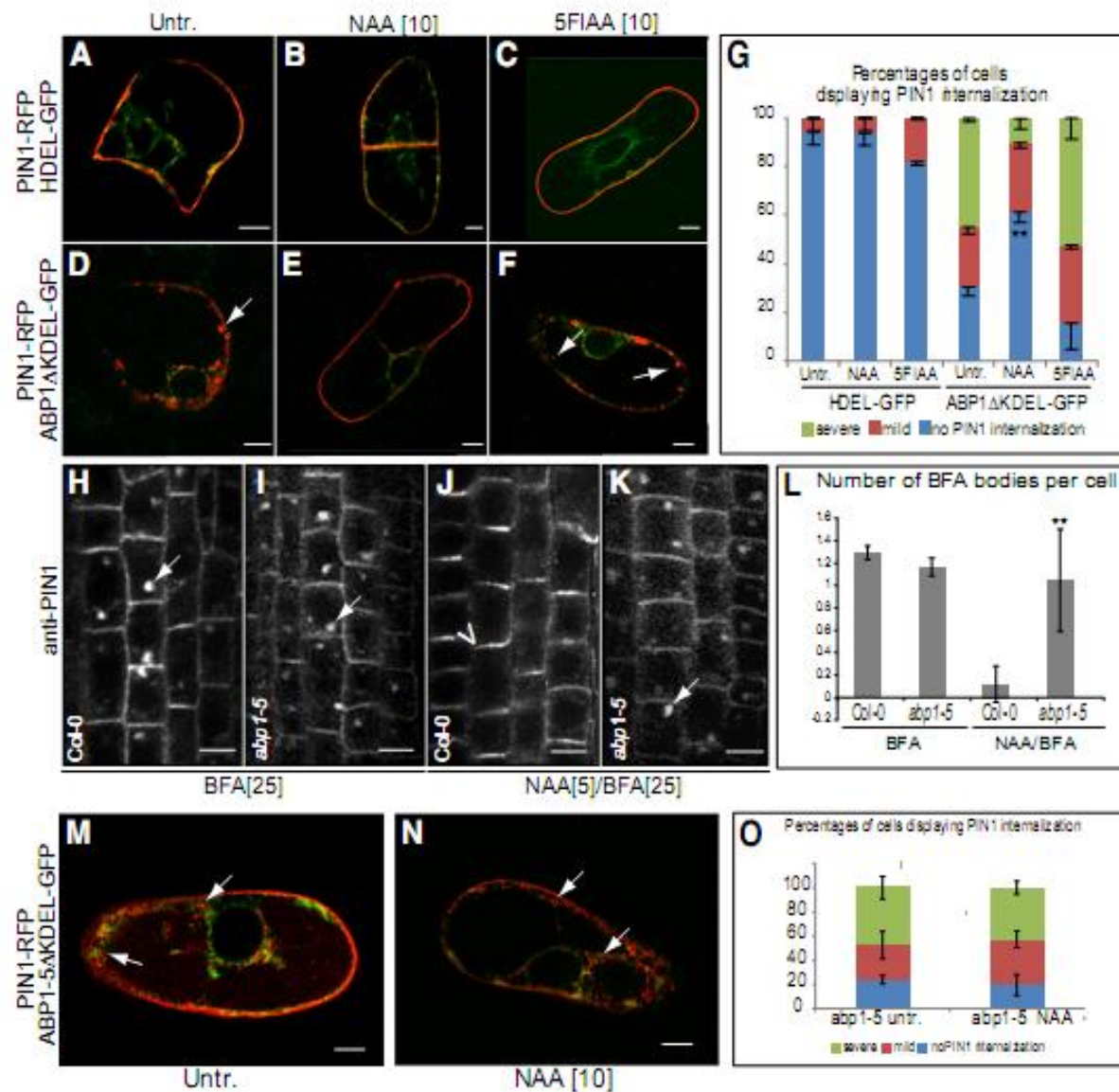
Auxin targets clathrin mechanism of endocytosis



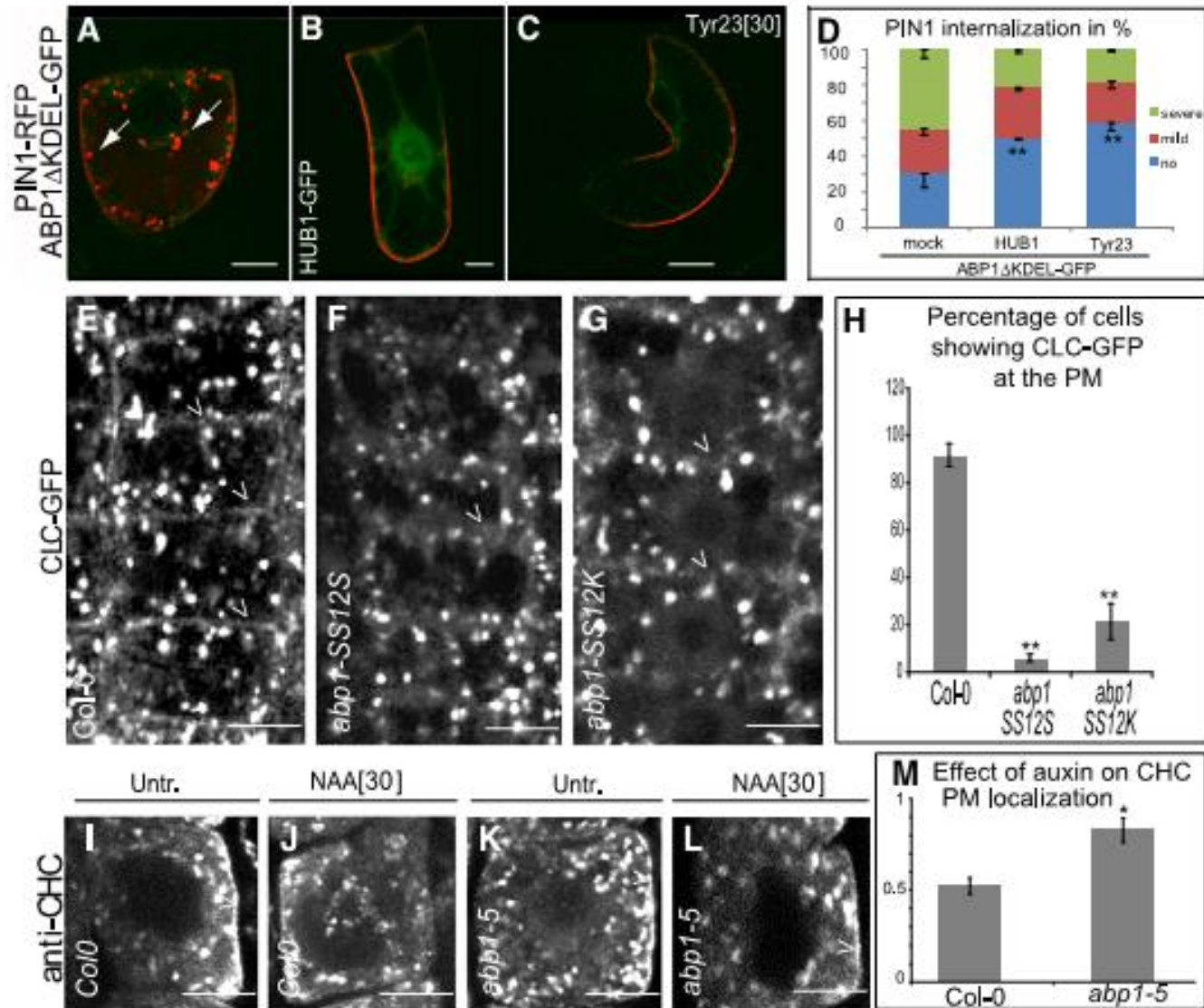
ABP1 positively regulates endocytosis



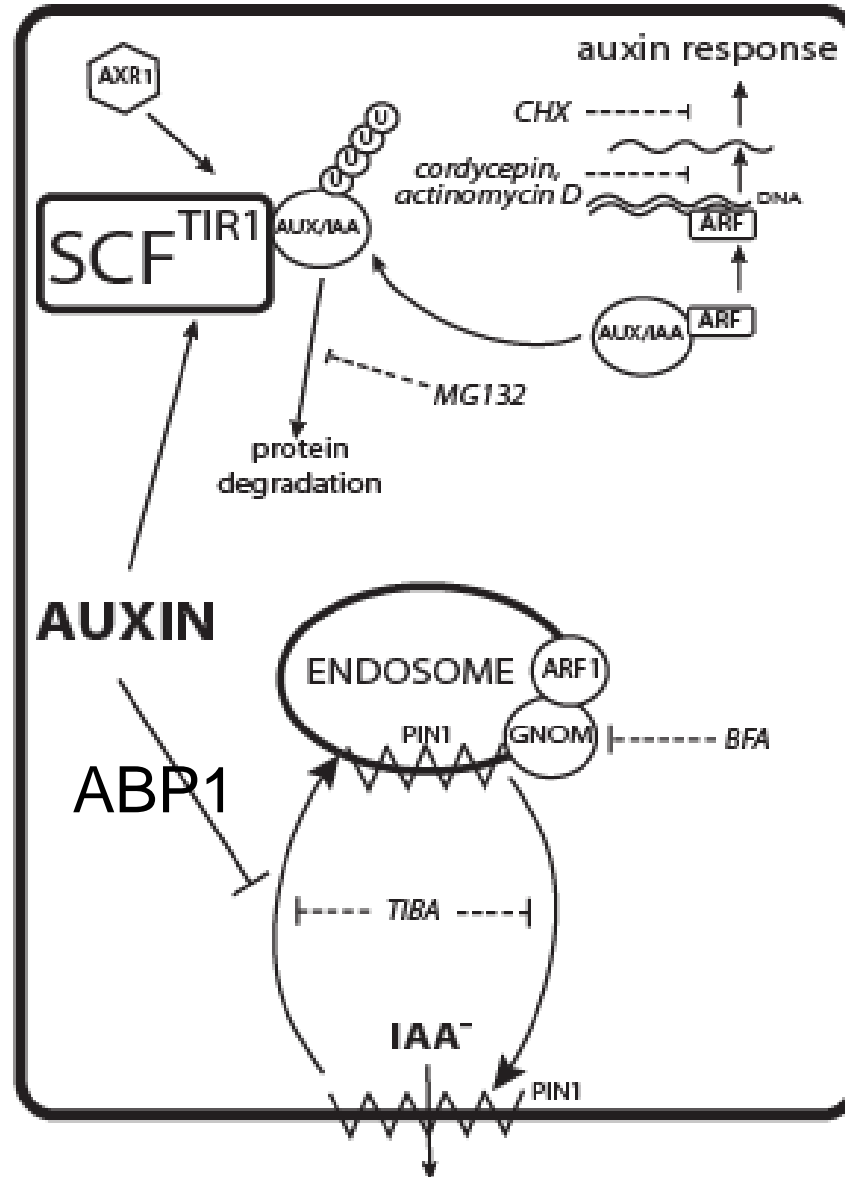
ABP1 mediates auxin effect on endocytosis



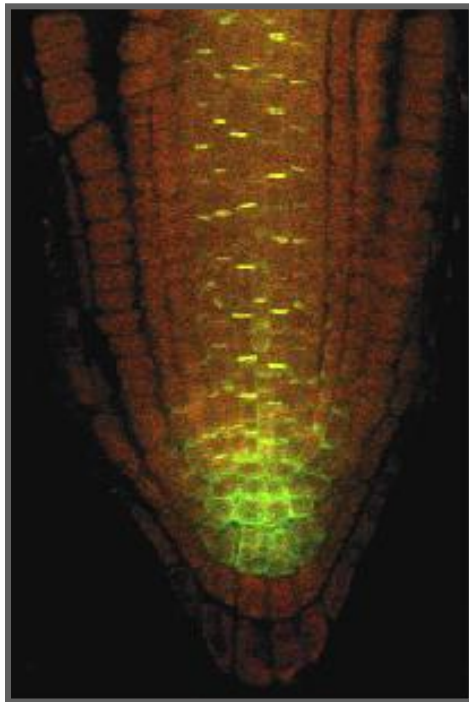
ABP1 mediates auxin effect on clathrin



ABP1- and TIR1-dependent Signaling



Mutant Screen for Components of PIN Polarity and Cycling



PIN:GFP

EMS mutagenesis.
Screening for
polarity and cycling
defects.



intragenic

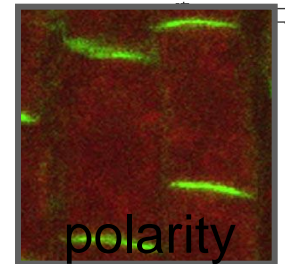
extragenic

sequencing

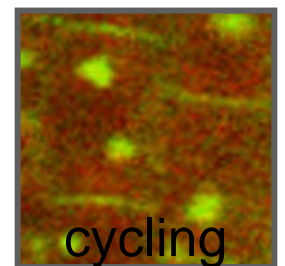
cloning

**important
residues**

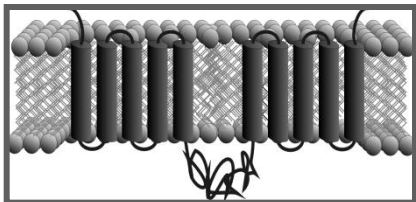
novel genes



polarity



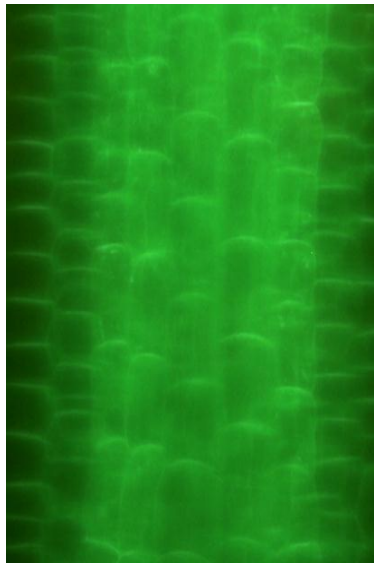
cycling



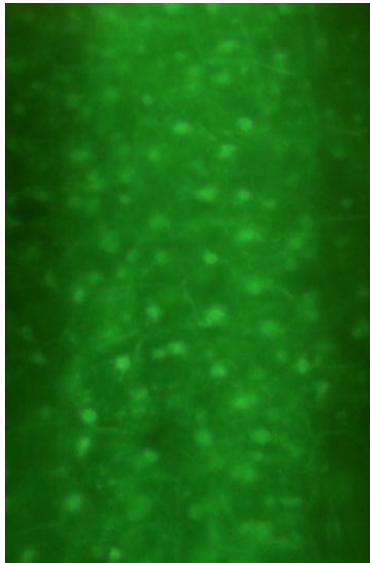
“Cell Biological” Mutant Screens in Progress:

Auxin effect on endocytosis: 3 confirmed mutants

30' NAA 30 μ M/90' BFA 50 μ M

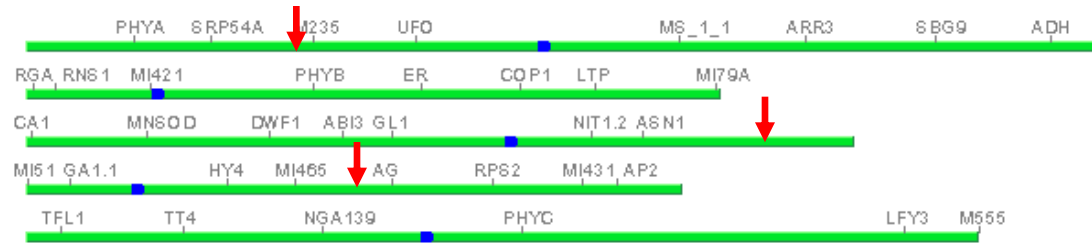


wt



mutant

Auxin-resistant BFA patches mutants



Novel Pathway for Auxin Signaling

Auxin inhibits endocytosis including internalization of PIN proteins

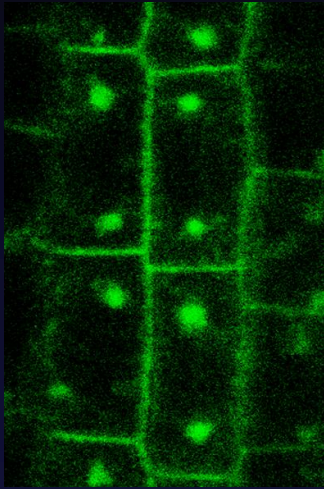
This is mechanism by which auxin stabilizes PINs at the cell surface thus stimulating auxin efflux.

This auxin effect involves novel, genetically tractable auxin pathway

Auxin Inhibits PIN Internalization and Stimulates its Transport

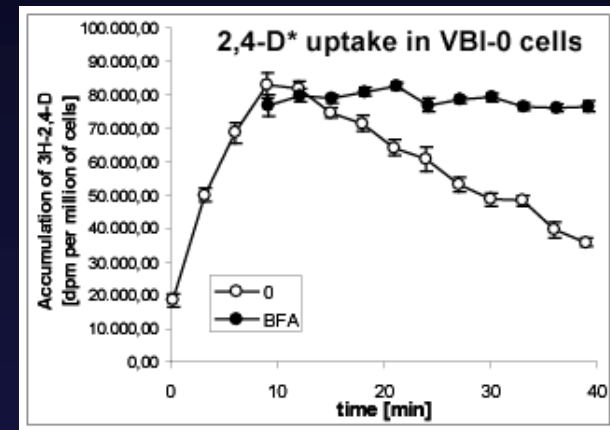
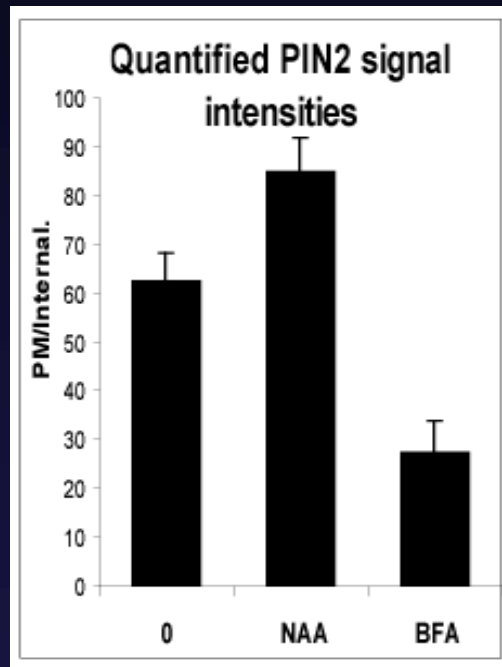


BFA



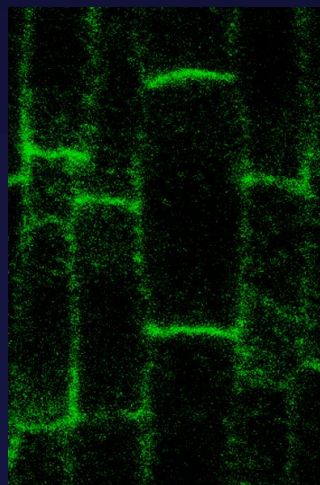
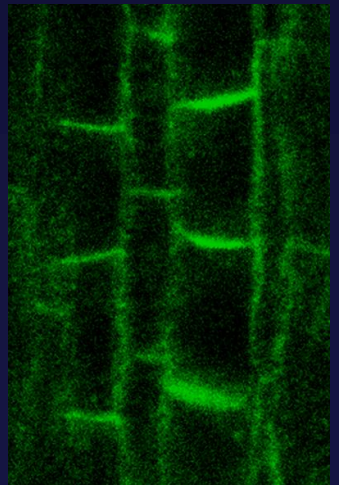
PIN2 at PM

Auxin efflux

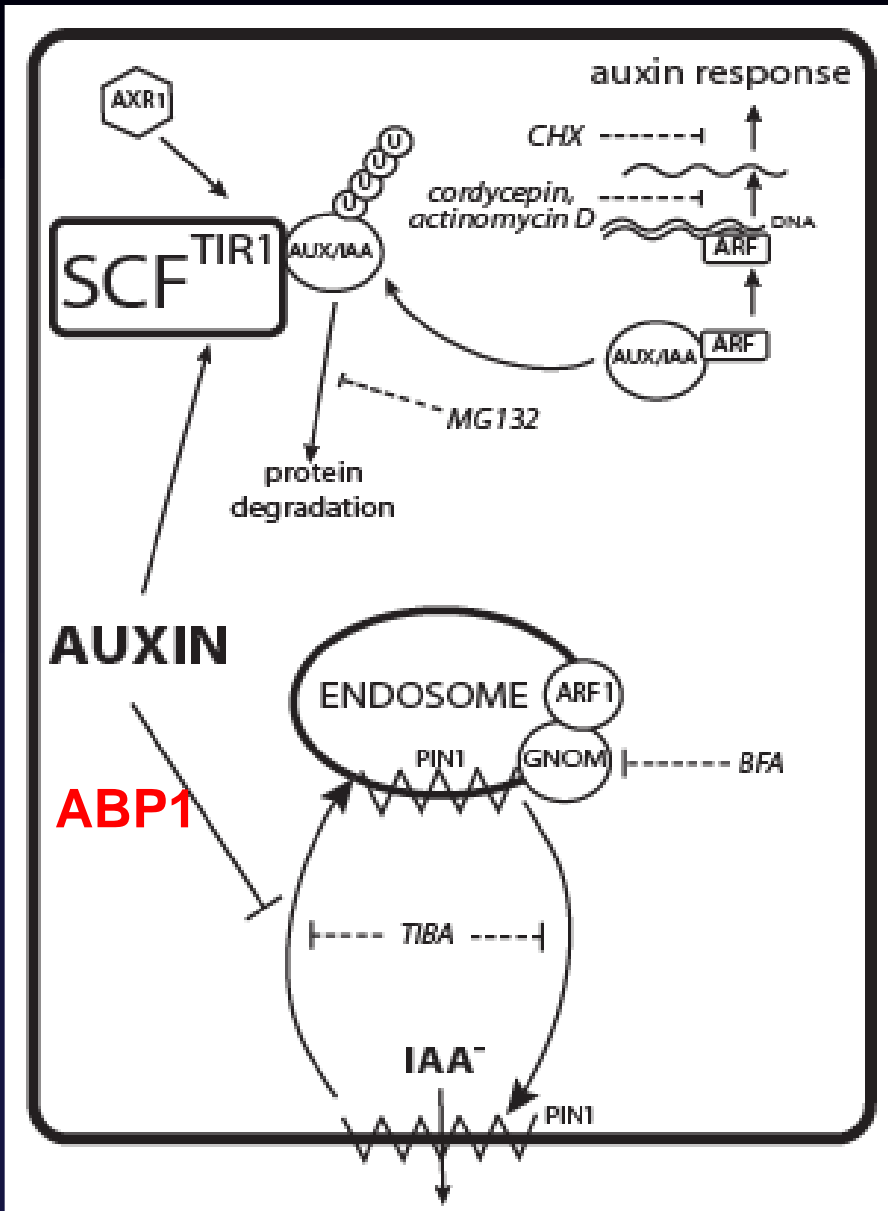


NAA/BFA

2,4-D/BFA

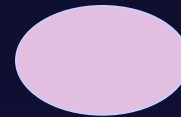


Auxin Signaling for Endocytosis

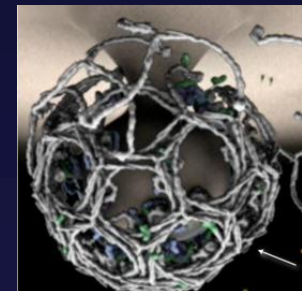
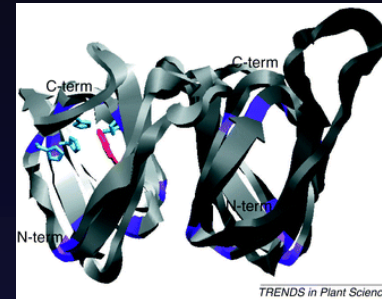


Auxin

ABP1

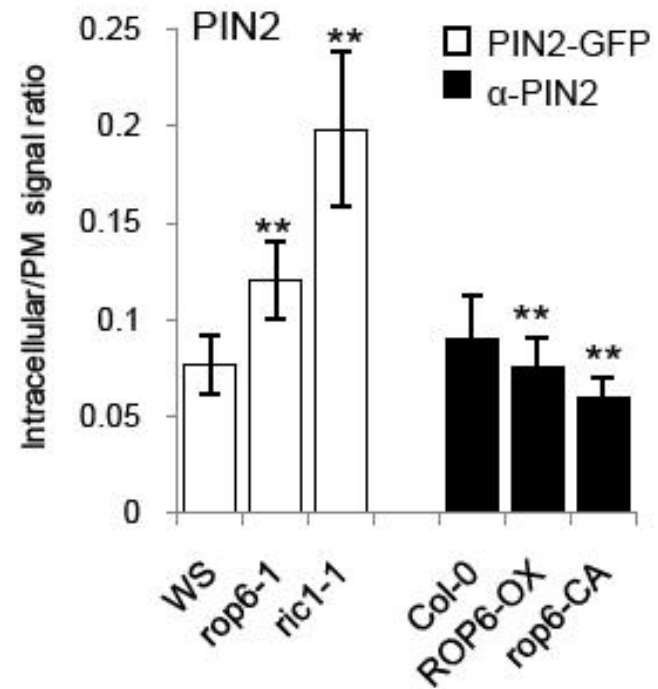
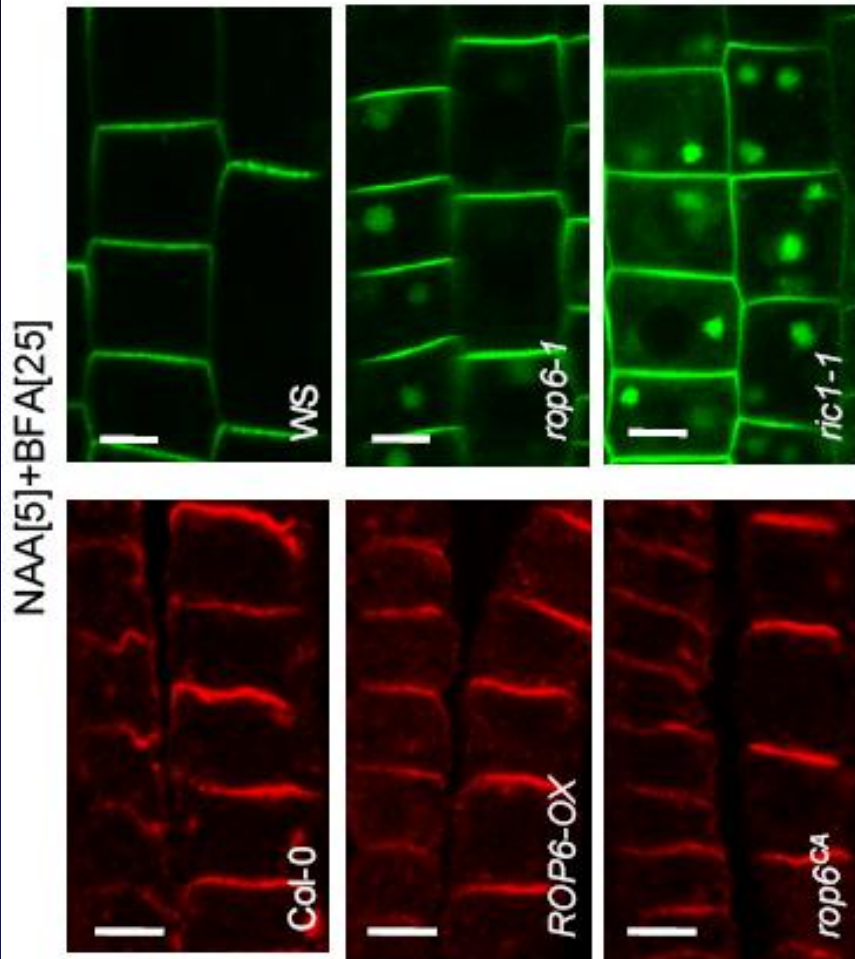
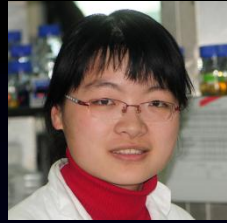


Clathrin

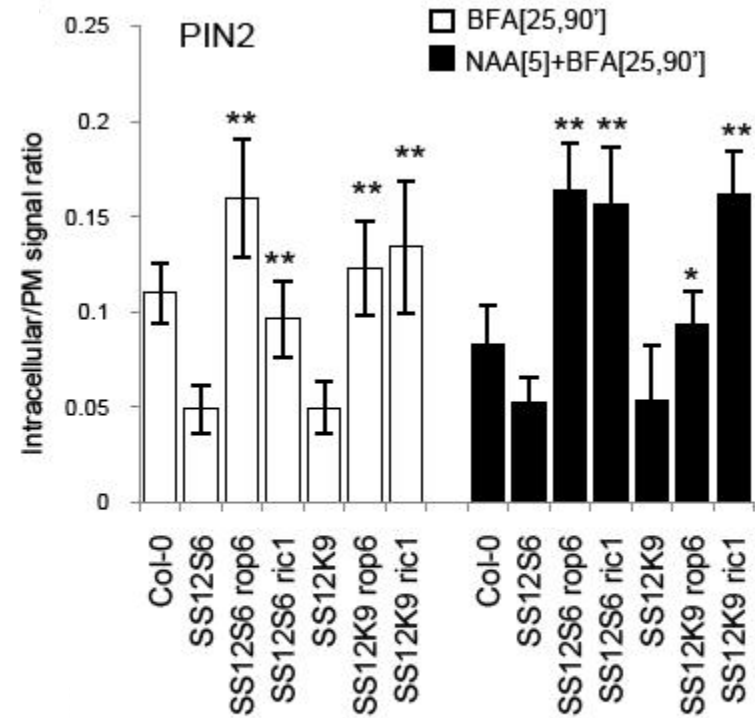
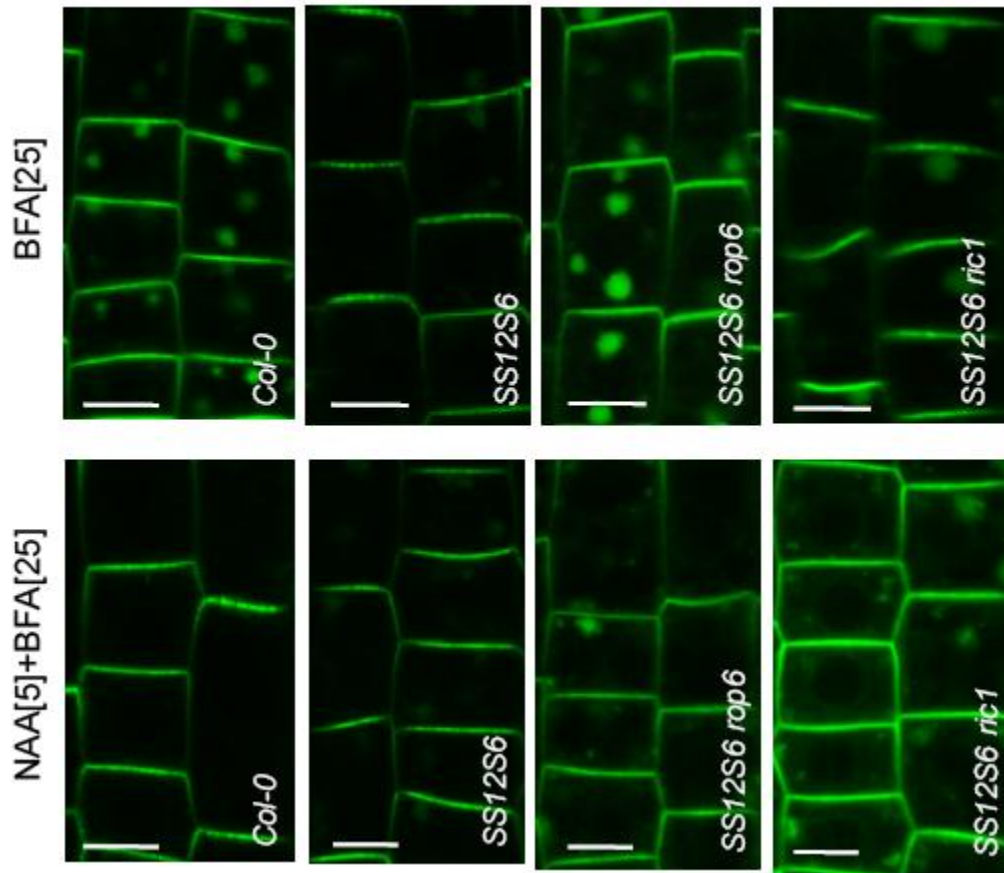
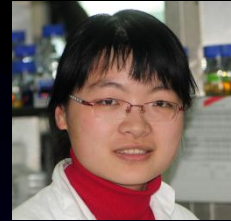


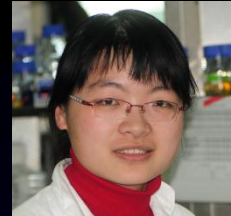
PIN endocytosis

Auxin effect on PIN endocytosis requires ROP6/RIC1 activity



ROP6/RIC1 acts downstream of ABP1 to regulate endocytosis





ABP1/ROP signaling for endocytosis and polarity

Auxin

ABP1

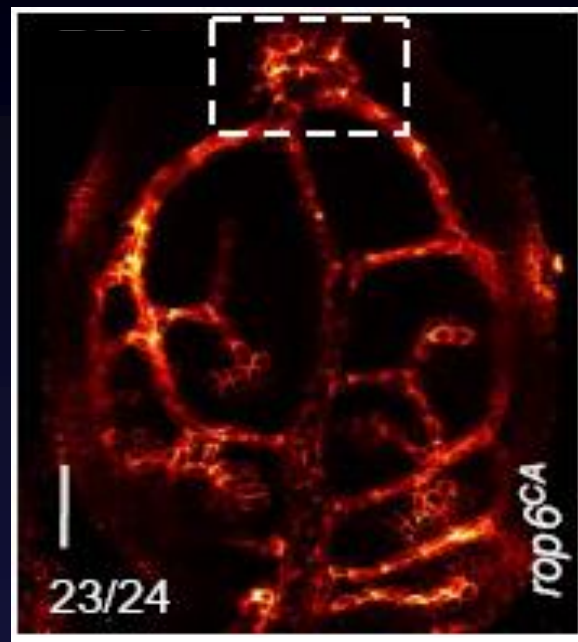
ROP

Clathrin

PIN endocytosis

PIN polarity

Vascular tissue formation



Gravitropism

