

# Explorations of Noise in the Shoot Apical Meristem

Henrik Ahl

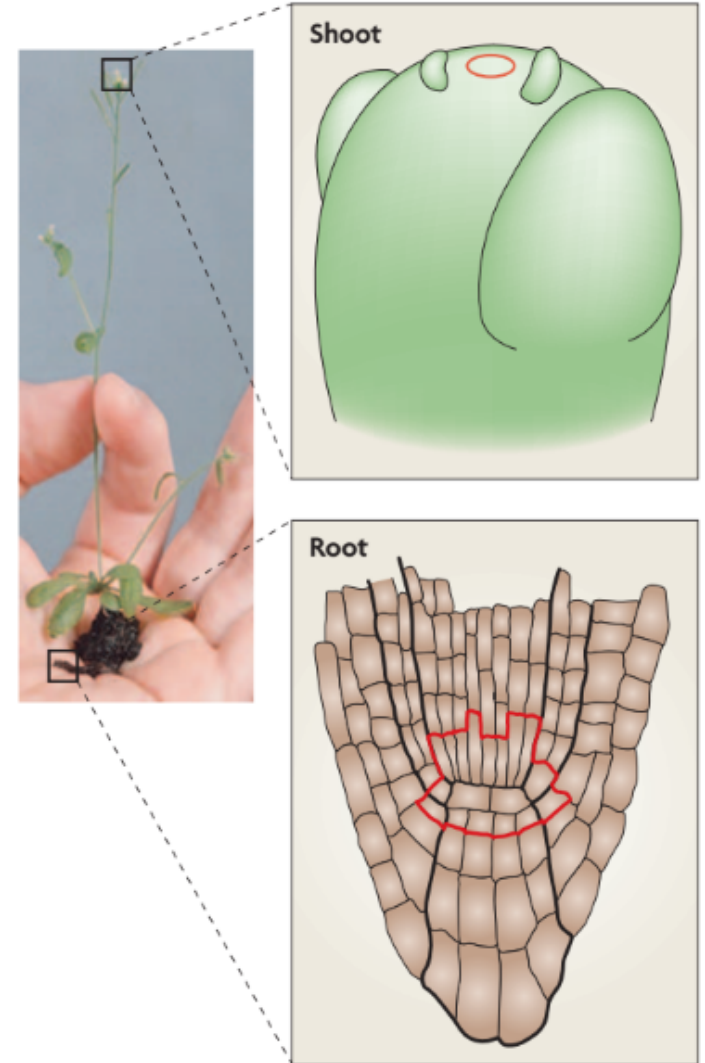
MPhil in Computational Biology

University of Cambridge

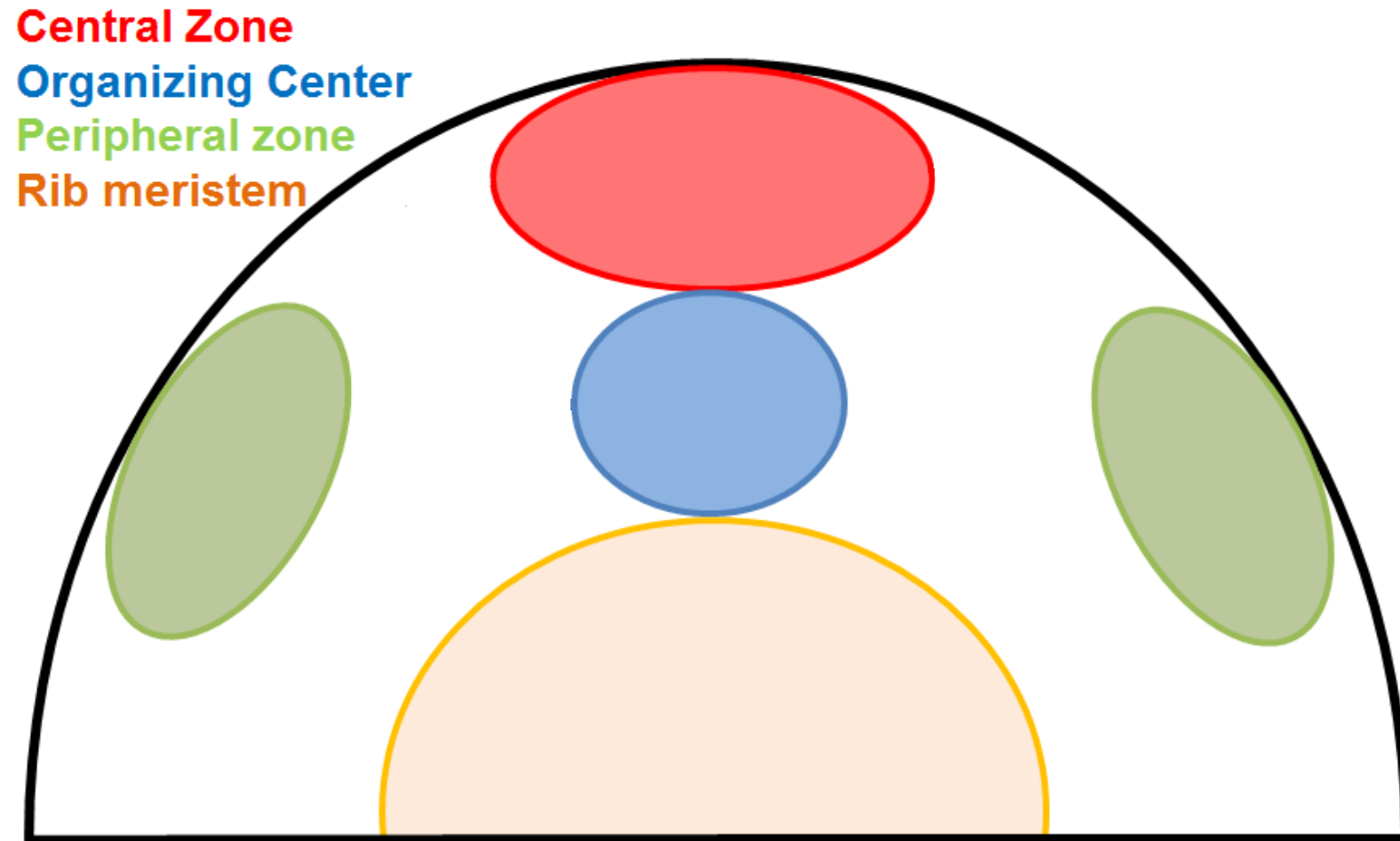


# The SAM is a developmental hub in *A. thaliana*

- Development of **aerial organs** (flowers, leaves, tissue) is determined at the SAM
- The stem cell population is located at the **apex**
- Gene expression specifies **cell identity**
- 
- 



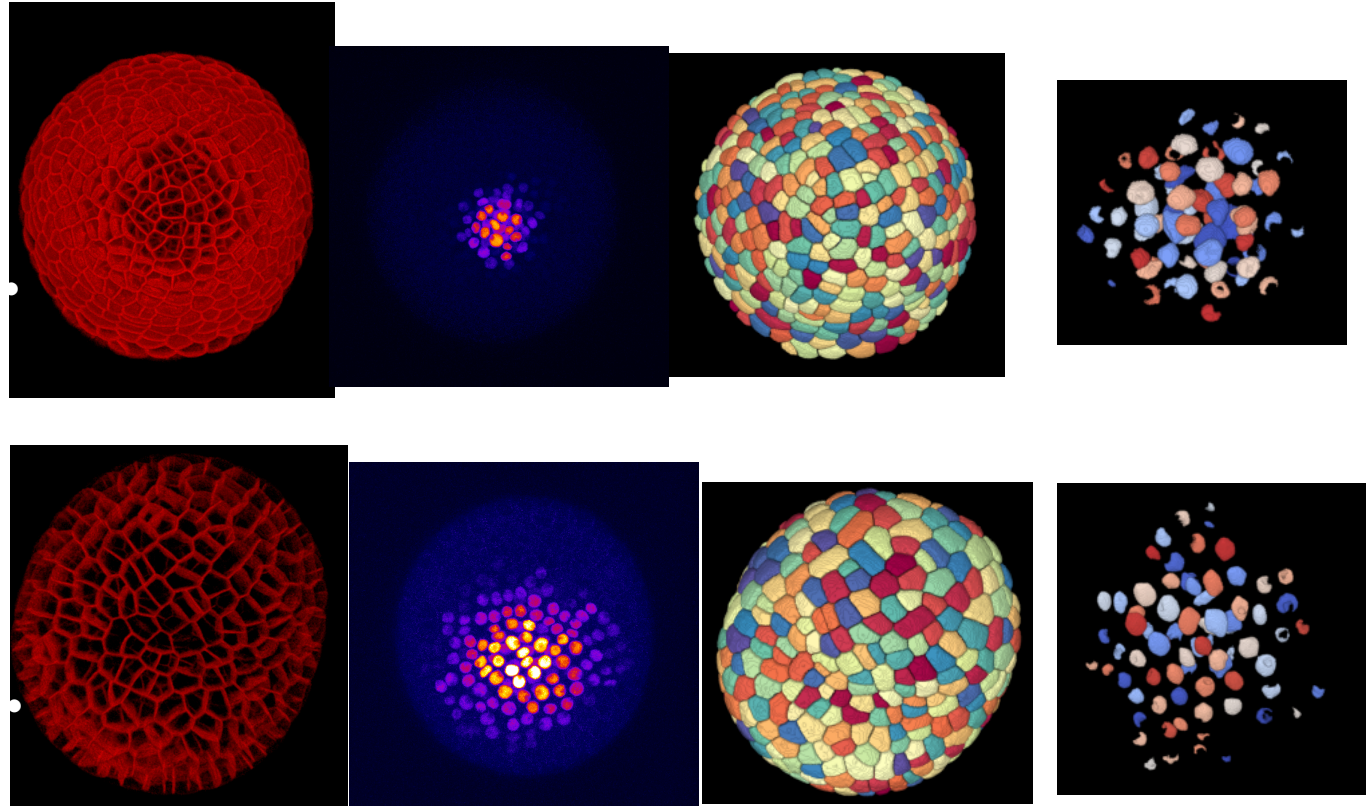
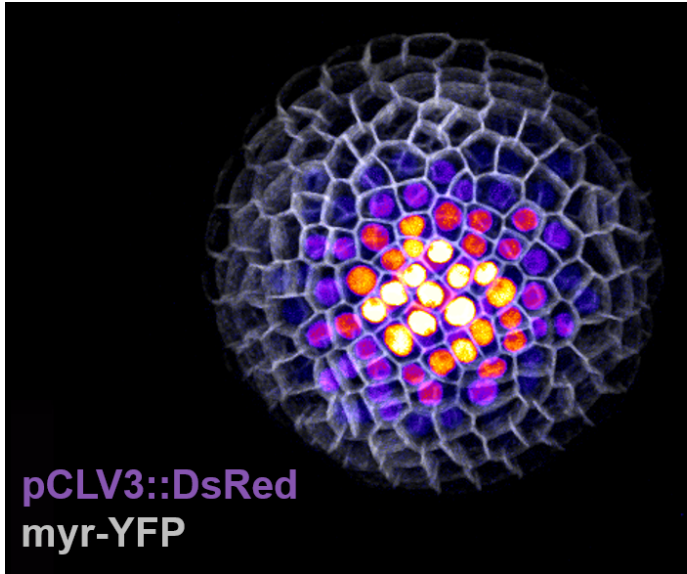
# negative feedback loops



# Project outline

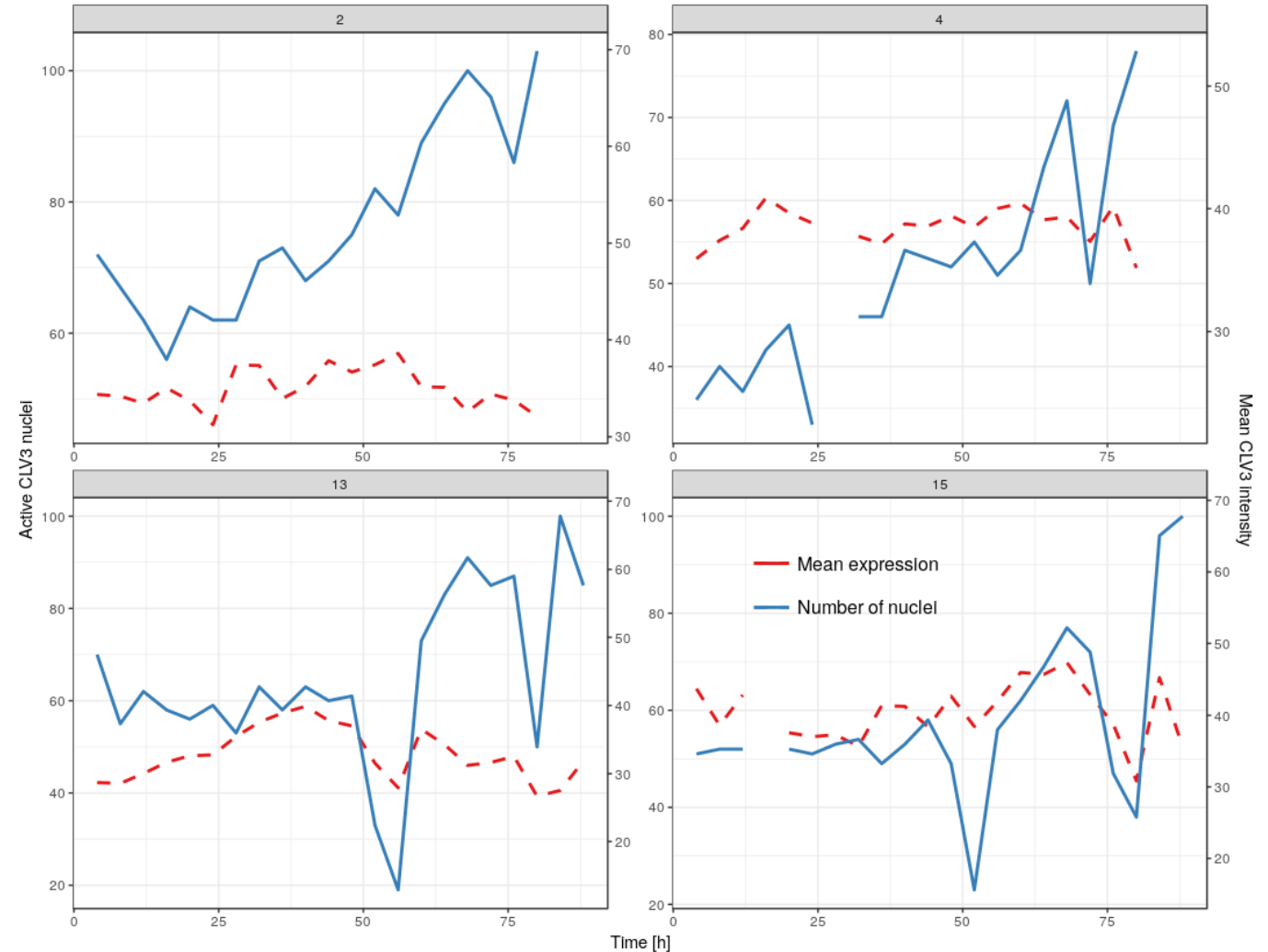
1. Design of data analysis pipeline:
  - Tidies, prunes and computes segmentation, cell line and division data
  - Written in R
  -
2. Exploration of data used in Willis et al. (2016)
3. Stochastic model investigations of data findings

# Data: High-resolution timecourses of plant growth



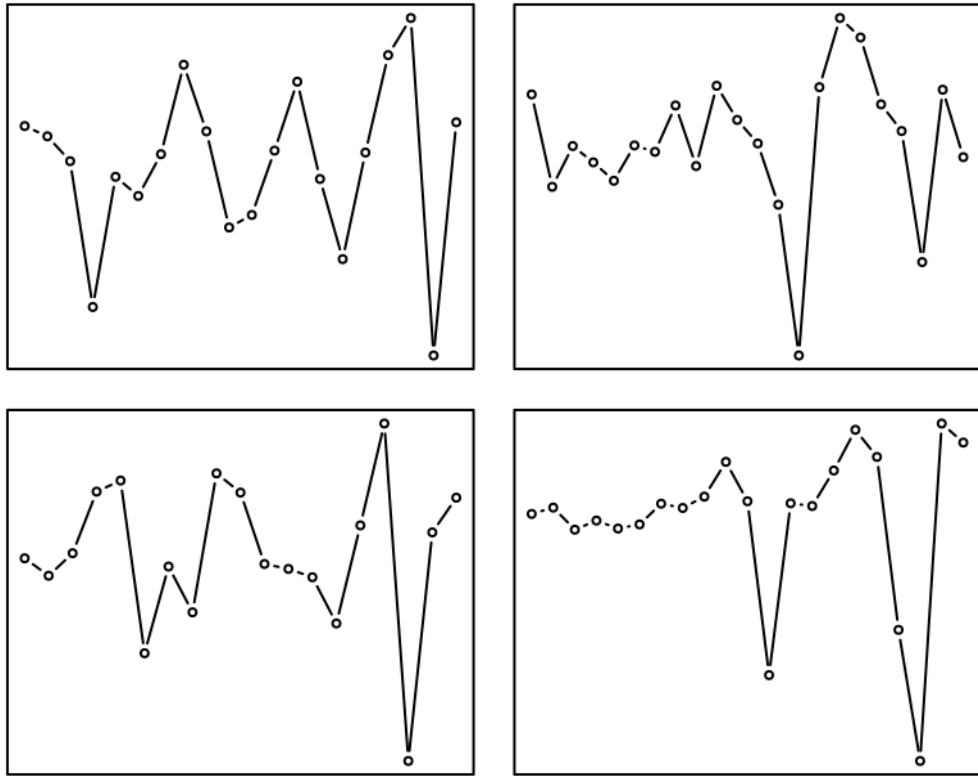
# Data shows large perturbations I

- Dilution of NPA perturbs system
- Oscillations **not linked** to mean expression
- Visually observable domain fluctuations in raw images



# Data shows large perturbations I

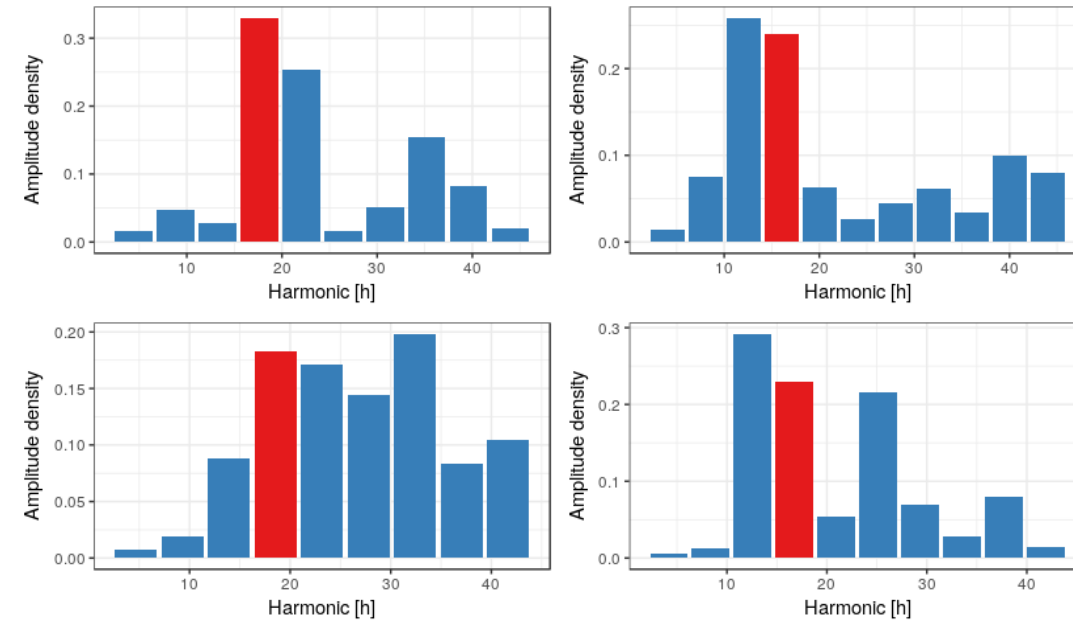
Detrended curves



Fourier  
decomposition

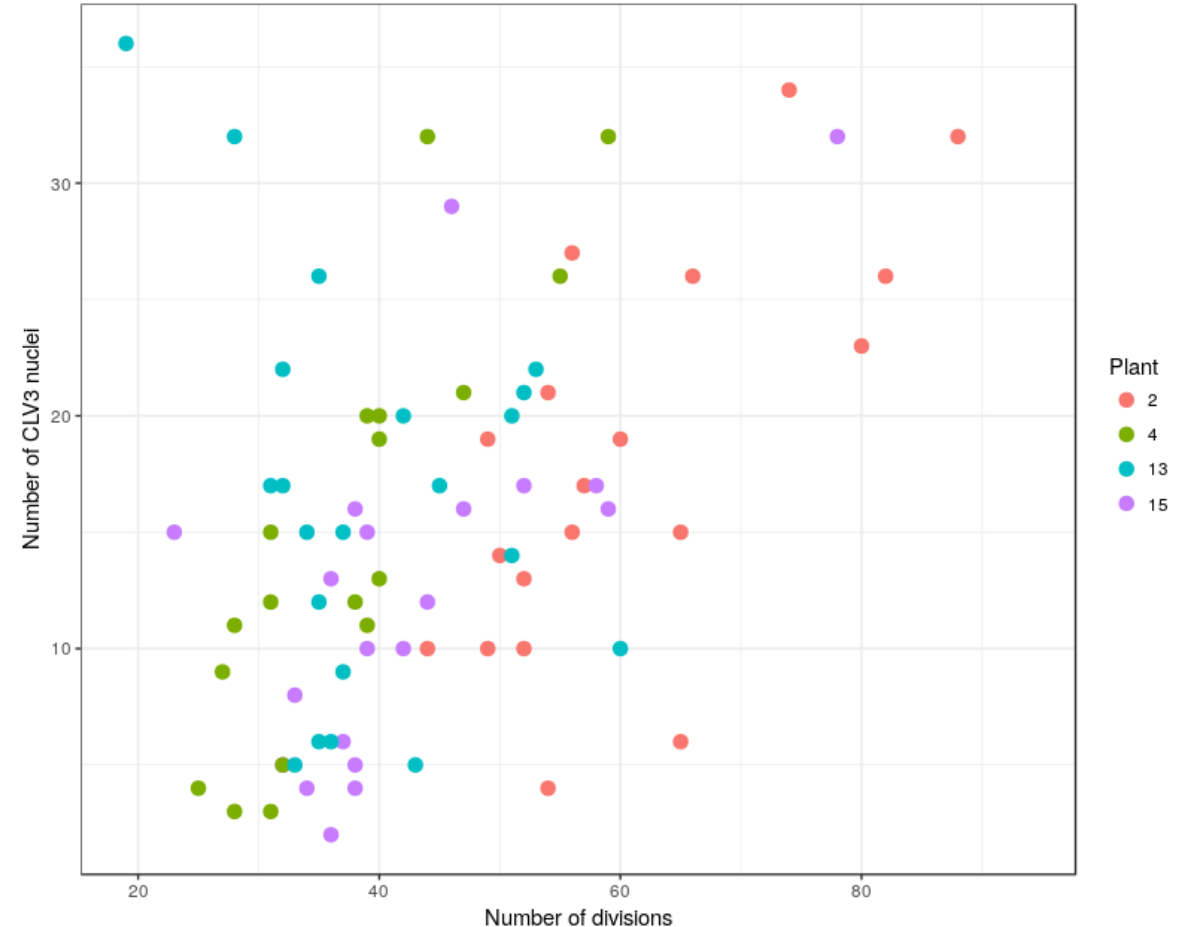


Frequency density



# Data shows large perturbations II

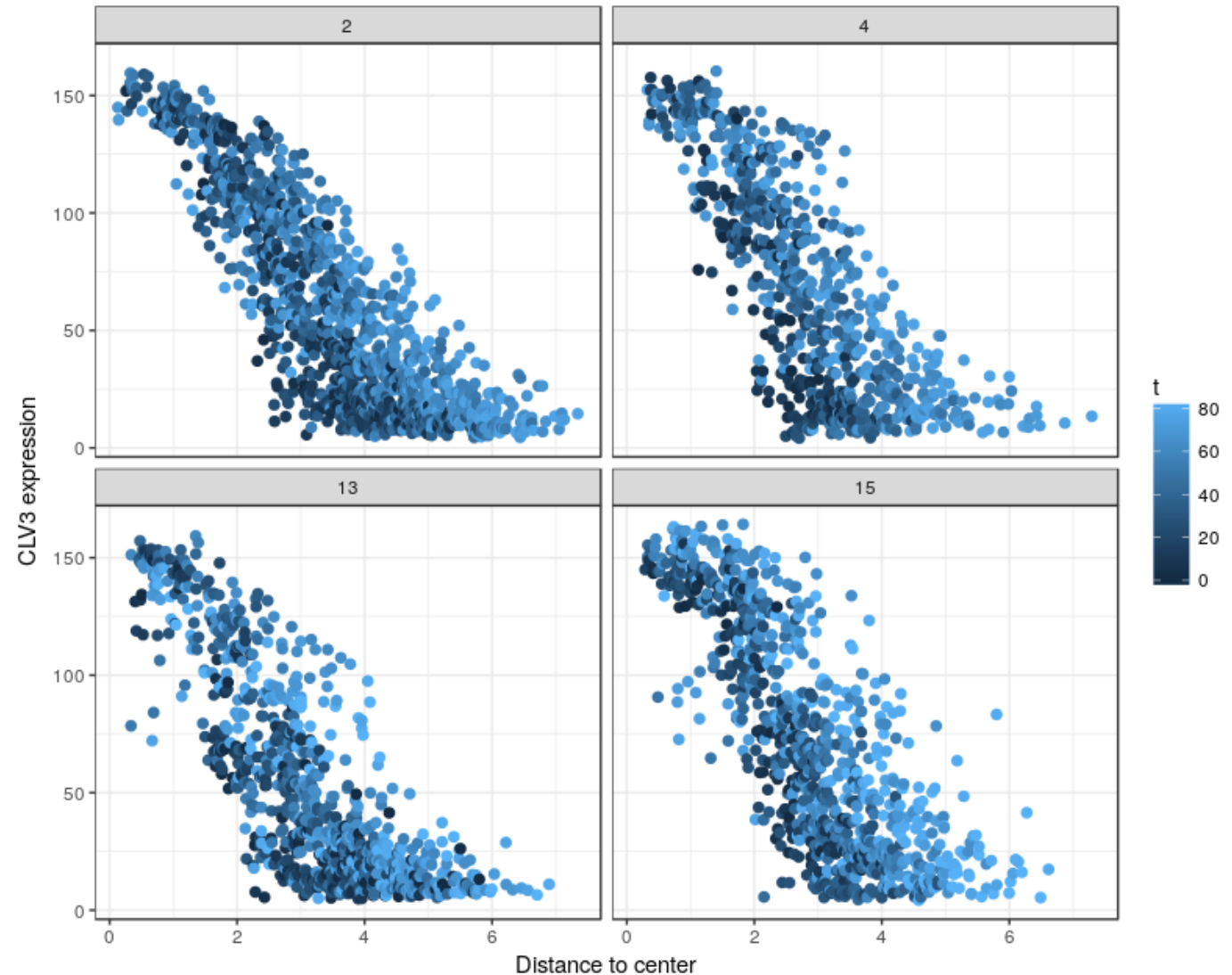
- #Nuclei correlates with #Divisions
- All:  $R = .43$ ,  $p = 6.01e-05$ 
  - Plant 2:  $1.90e-04$
  - Plant 4:  $6.79e-06$
  - Plant 13:  $0.21$
  - Plant 15:  $8.38e-05$
- Plants appear to undergo significant **functional** variance



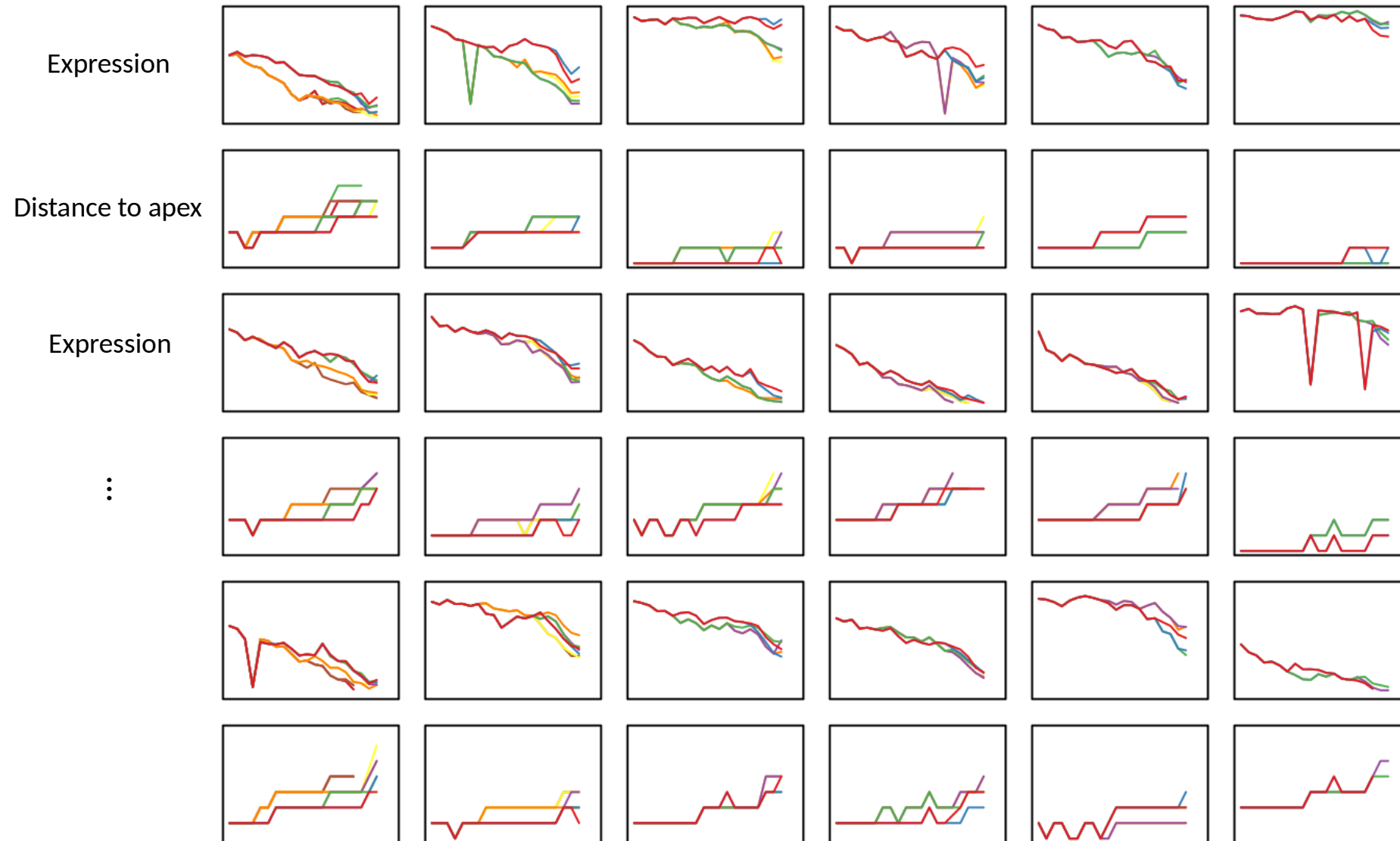


# Cells at apex behave robustly I

- Analysis hampered by definition of apex
- Drops in expression at apex likely from **loss of signal** at division
- Investigations of bimodality show **no results**



# Cells at apex behave robustly II



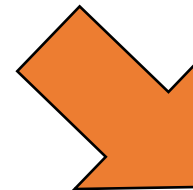
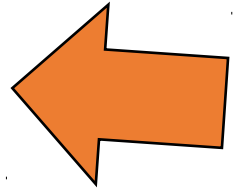
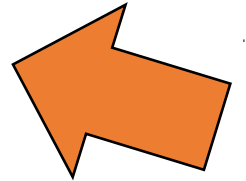
# Cells at apex behave robustly II

Expression

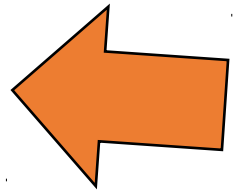
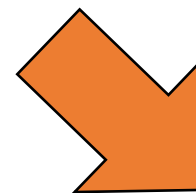
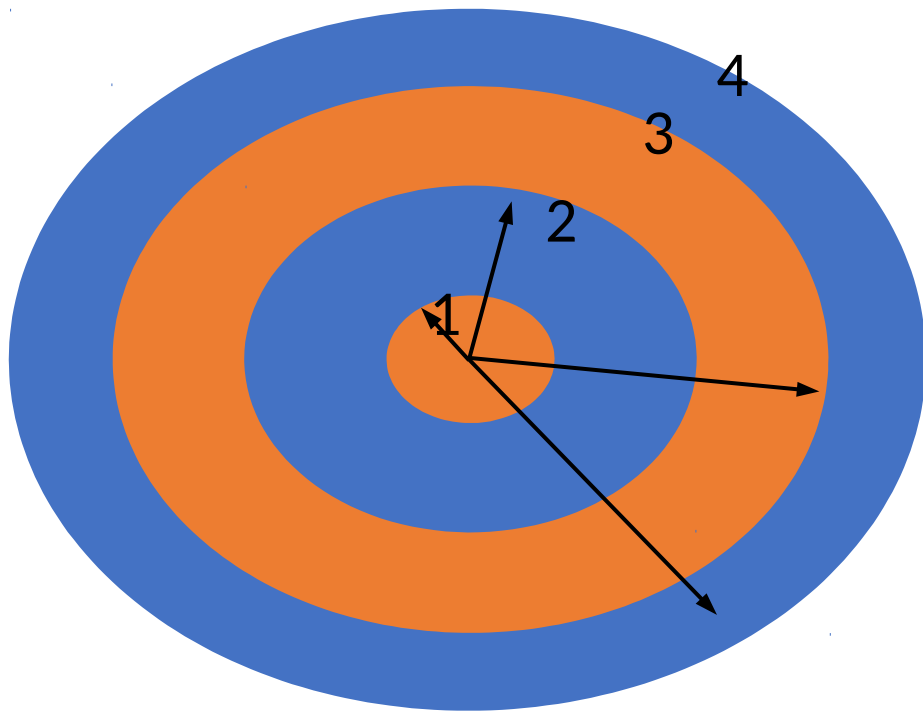
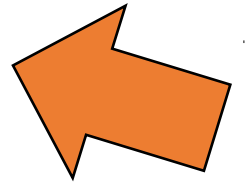
Distance to apex

Expression

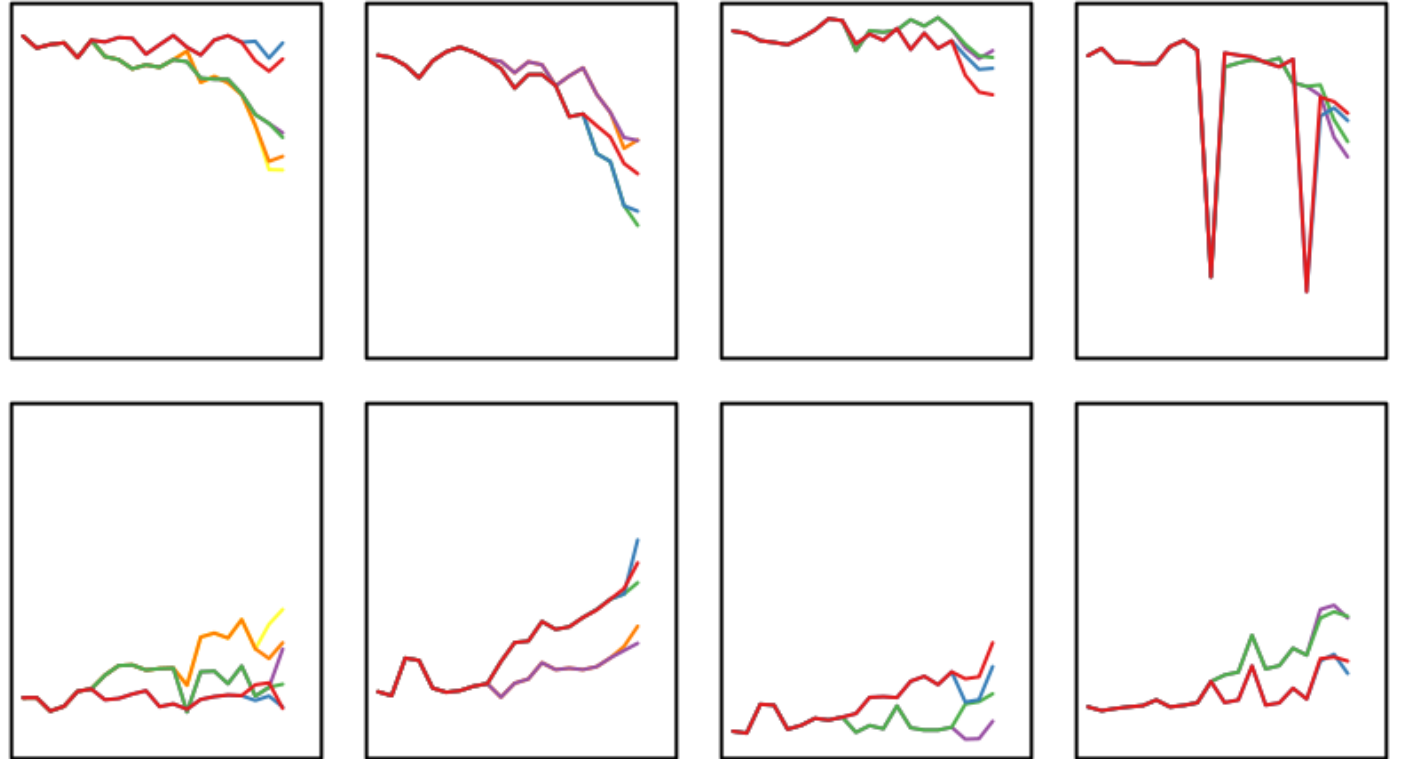
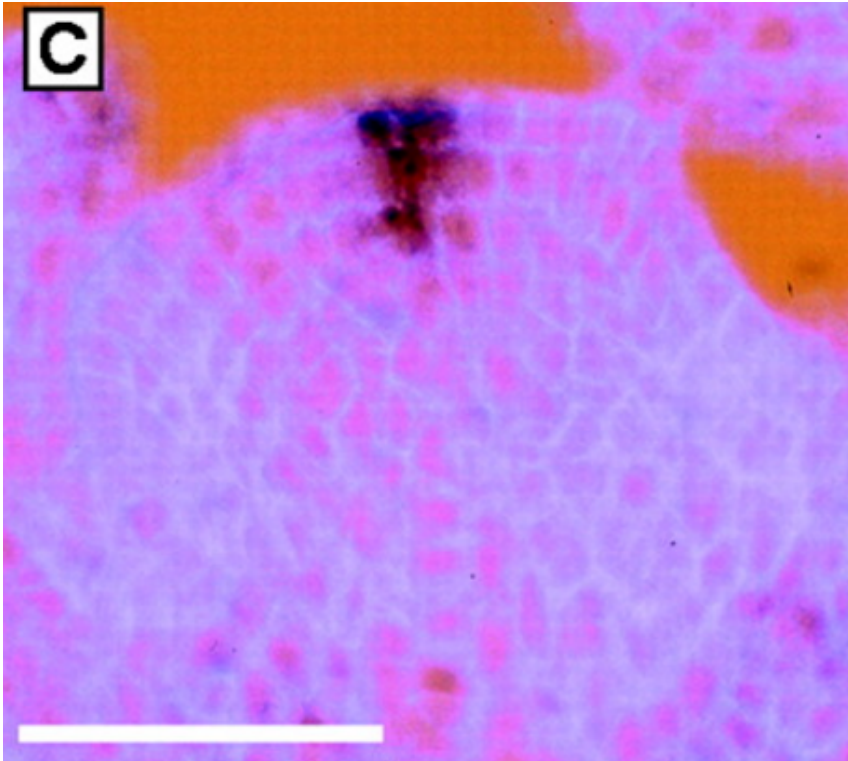
⋮



# Cells at apex behave robustly II



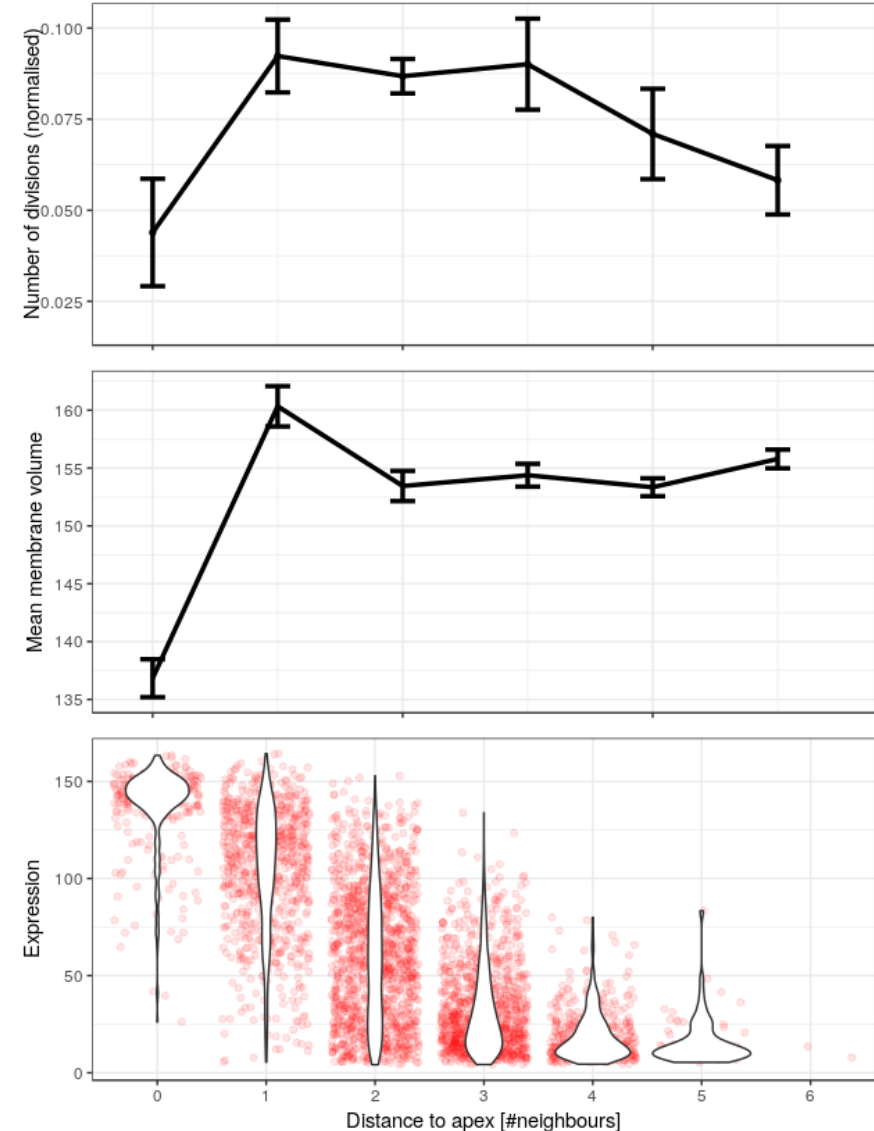
# Cells at apex behave robustly II



Fletcher et al. (1999)

# ... and just differently

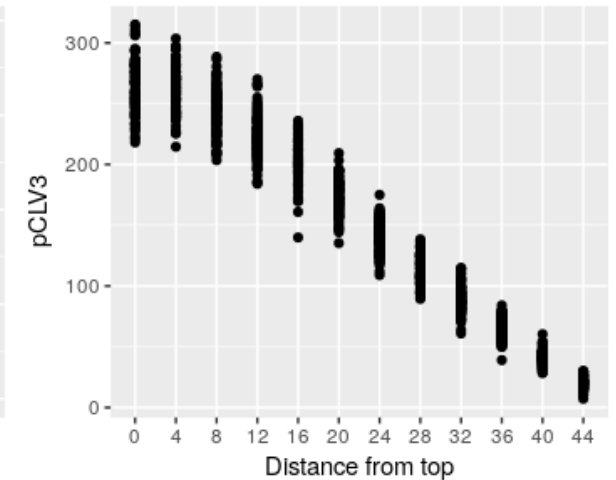
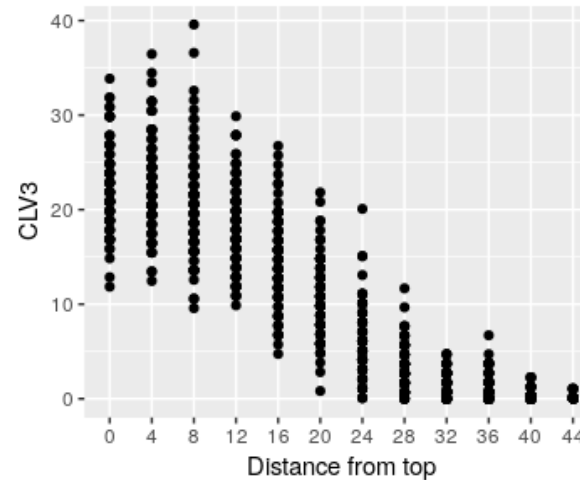
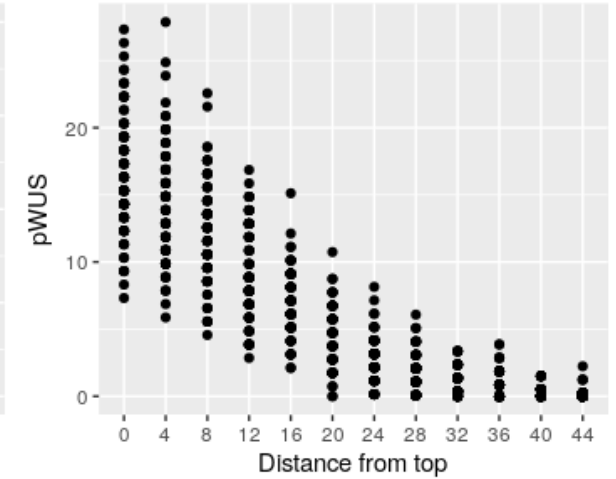
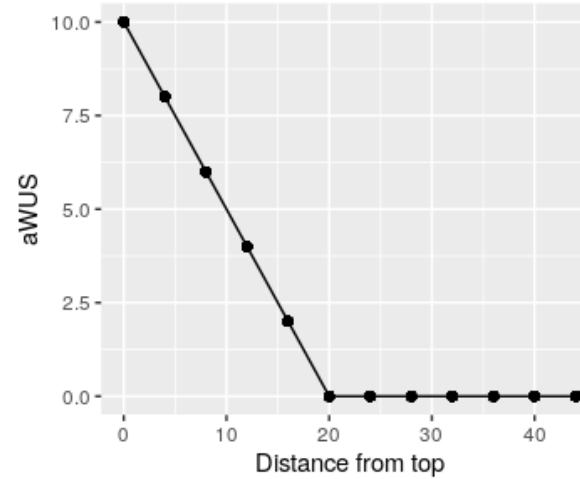
- Top cells divide **less frequently** due to a lower growth rate
- Average membrane volume at apex **significantly lower** than in other periclinal layers
- 
- CLV3 expression **not related**
- 



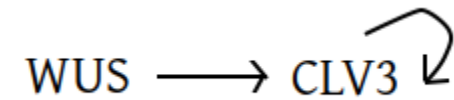
# A simple model tests the possible regulation

WUS  $\longrightarrow$  CLV3

- Epidermal 1D model
- Gillespie and / or Milstein's method
- Optimised using genetic algorithms
- Cannot suppress noise at apex



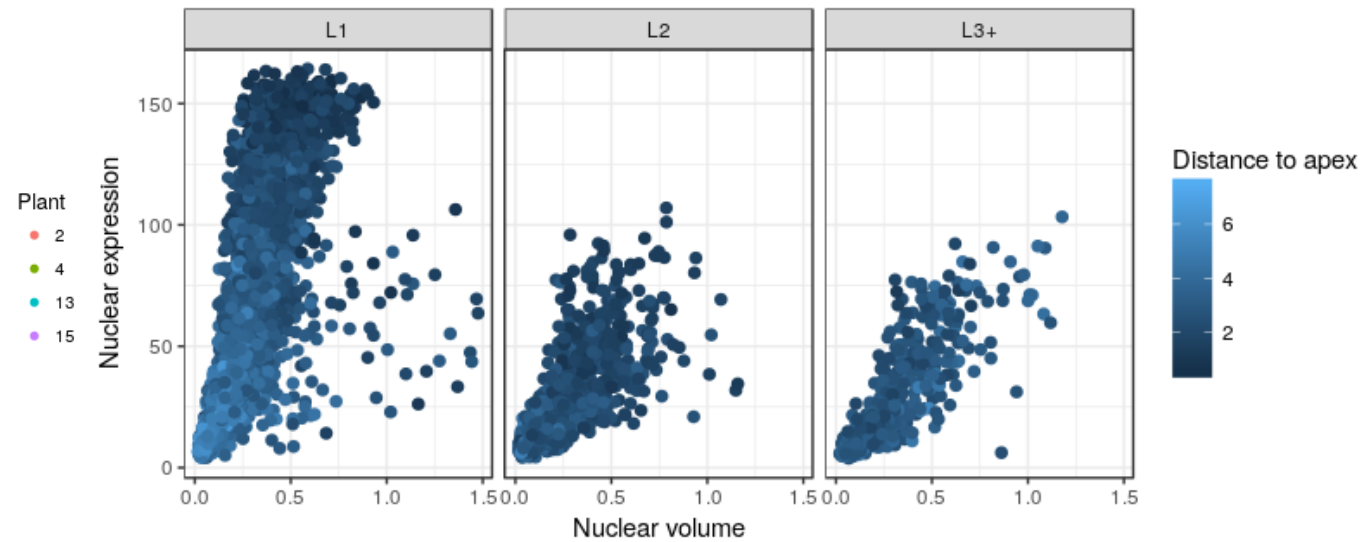
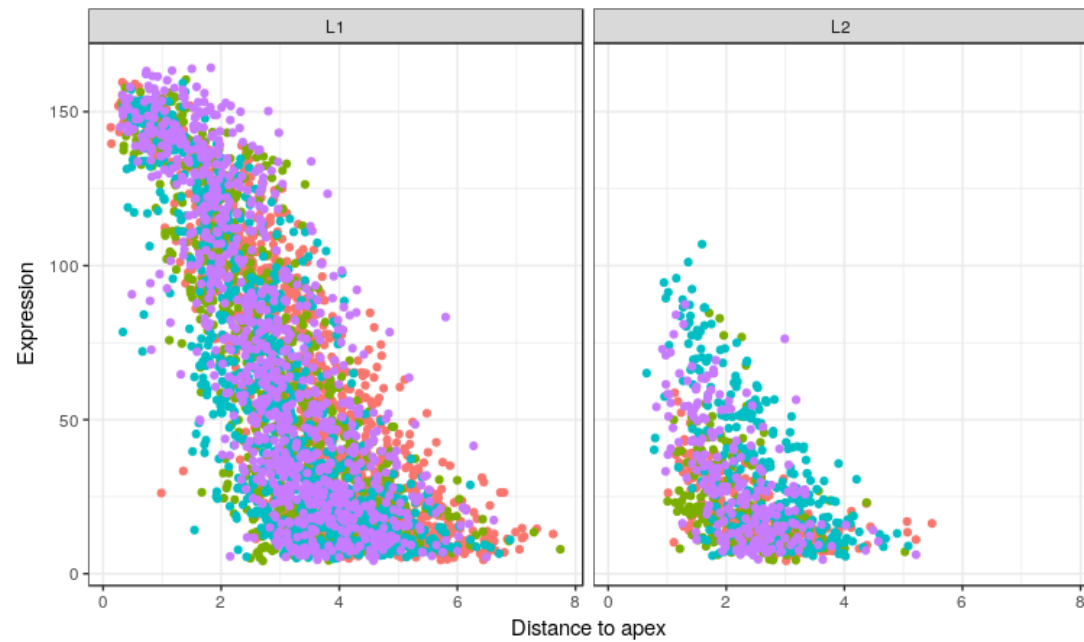
# A simple model tests the possible regulation



- Possibly? More work needed.



# Layer analysis suggests L1 regulation



# Acknowledgements

José

Henrik

Jérémy

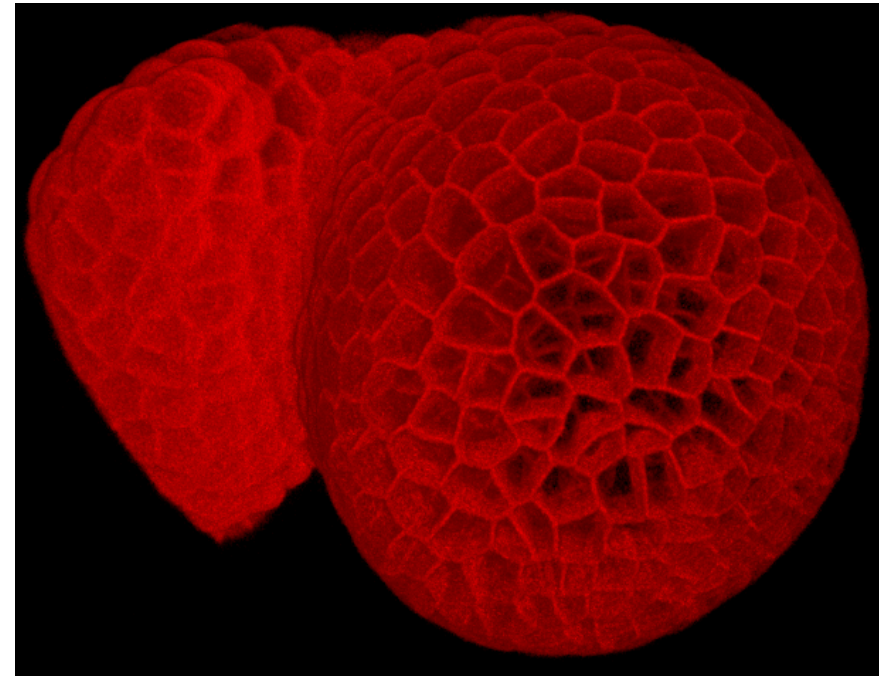
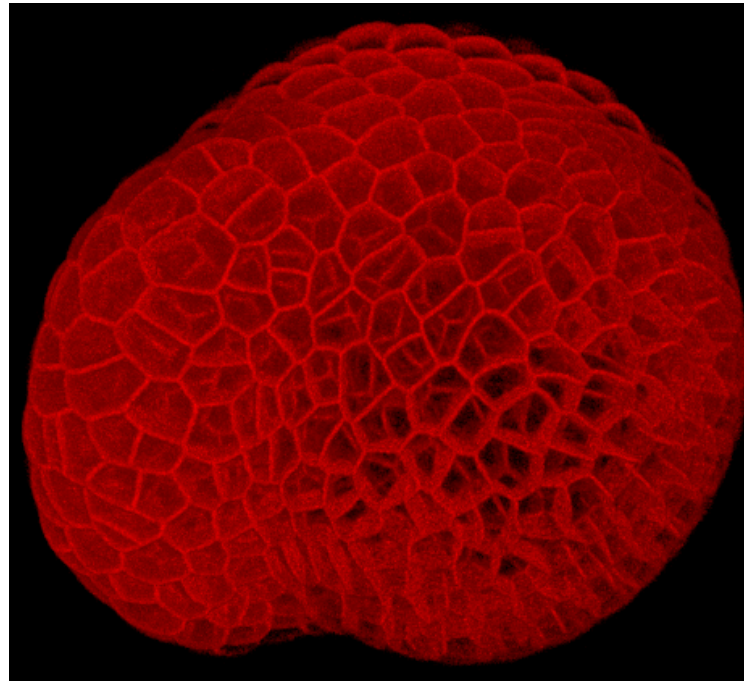
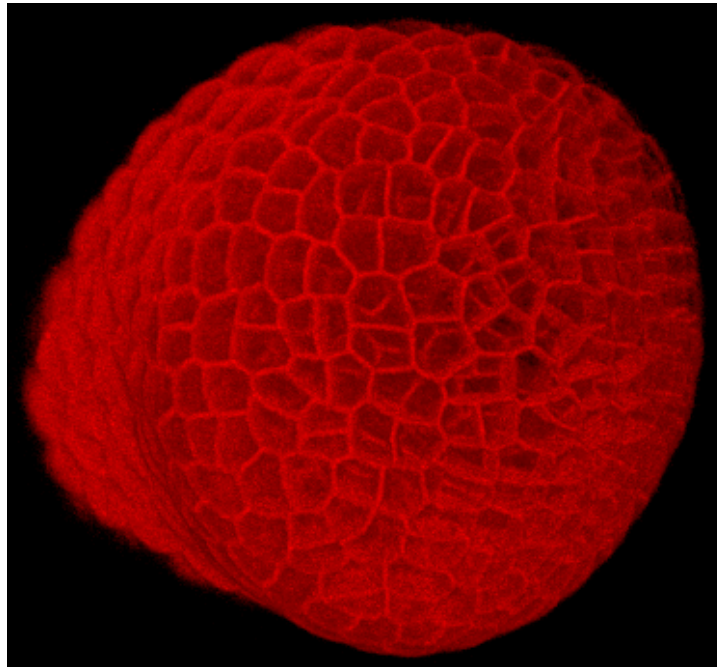
Yassin

+ everyone else

Thank you!



# Initiation of organs implies NPA dilution



# Additional results: Longevity

- WUS domain affects  
longevity?
- Currently investigated

