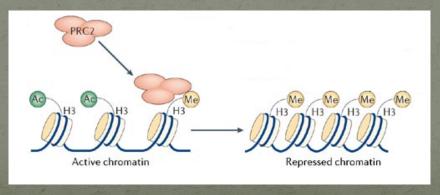
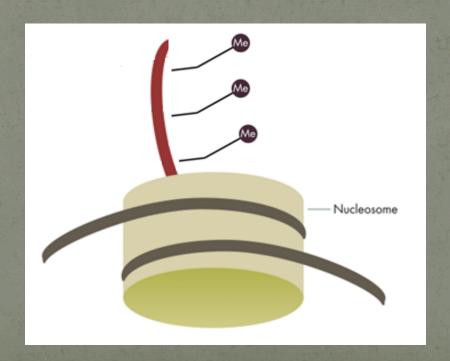
Identification of the Polycomb Repressive Complex 2 in Fungi

Phuong Pham Dr. Michael Freitag Summer 2012



Activating and silencing of genes

- Transcriptional "gene silencing"
- Genes can be repressed by methyl groups on histones

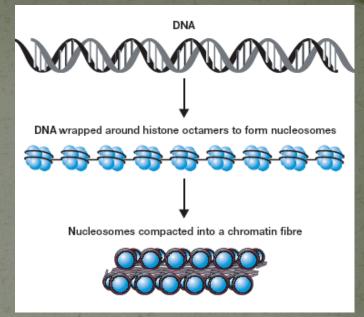


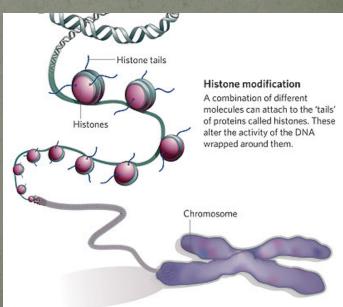
Function of gene silencing

- Regulate genes expression
- Normal development in stem cells
- Protect against viruses and transposons

Histone methylation

- Histones bind to DNA
- Makes up "chromatin"
- Histone 3:
 - methyl-lysine 4 (K4me2): active
 - methyl-lysine 27 (K27me3): silent





Organism

- H3K27me3 first found in fly
- Fusarium graminearum
- PRESENT: gene expression OFF
- ABSENT: gene expression ON
- Primary & secondary metabolites



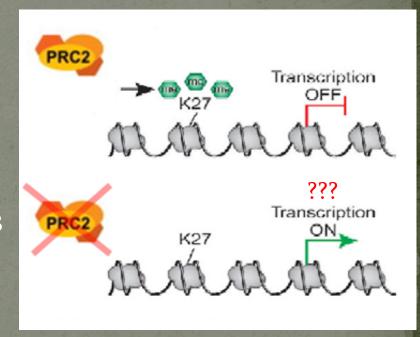


http://en.wikipedia.org/wiki/File:F.graminearum.JPG http://www.ag.ndsu.edu/pubs/plantsci/smgrains/pp894w.htm

Enzyme Responsible for H3K27me3

- Polycomb Repressive Complex 2 (PRC2)
- PRC2 includes 4 protein subunits:
 - Set7, Caf1-3, Suz12, Eed
 - (Subunits 1, 2, 3, 4)
- Deletion of subunit 1 removes H3K27me3
 - → infertility, slow growing, orange
- Developmental effects due to

misregulation



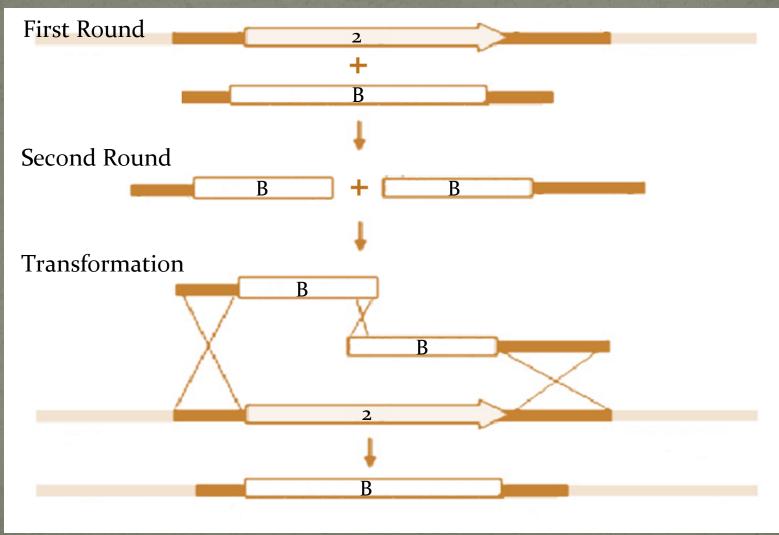
Aims of this study

- Test if deletion of subunits 2, 3, and 4 will show similar effects as deletion of subunit 1
- Study interactions between PRC2 subunits
- Determine the composition of fungal PRC2

Deletion by Split Marker PCR

- First round: Generated fragments
 - Caf1-3, Suz12, Eed (2-4)
 - *neo* and *hph* genes encoding for antibiotic resistance (A and B)
- Second round: Generated split marker fragments (flank and ½ of A or B)
- Transformation: Replaced genomic 2, 3, or 4 with A or B
 - Δ_2 (Δ_2 ::A and Δ_2 ::B)
 - Δ_3 (Δ_3 ::A and Δ_3 ::B)
 - Δ 4 (Δ 4::A and Δ 4::B)
- Only proceeded with Δ_2 ::A, Δ_3 ::B, and Δ_4 ::B

Split Marker PCR



Methods: Analysis of deletion strains

PCR and Transformation 1 and 1 a Transformant ChIP Analysis Grow Tissues RNA Sequencing Southern Western blotting blotting

Phenotype Check

PCR Check

Phenotype Check

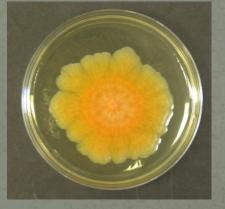


Wild Type (WT)

 Δ_1



Wild Type (WT)



 Δ 2

???



 Δ_3



 Δ_4

Methods: Analysis of deletion strains

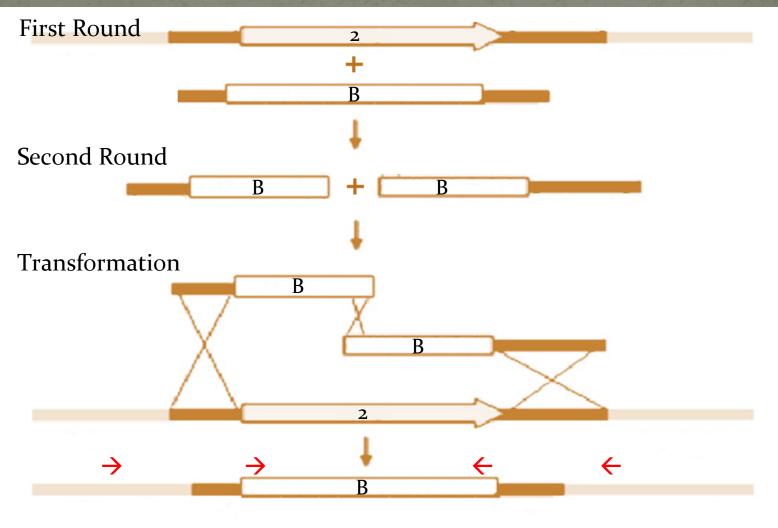
Phenotype

Check

PCR Check

PCR and Transformation 1 and 1 a Transformant ChIP Analysis Grow Tissues RNA Sequencing Southern Western blotting blotting

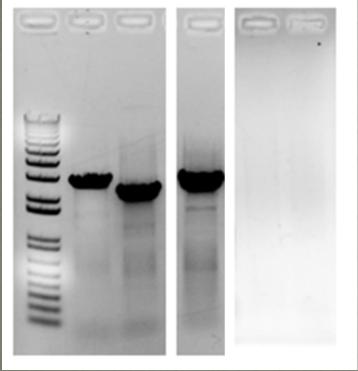
Split Marker PCR



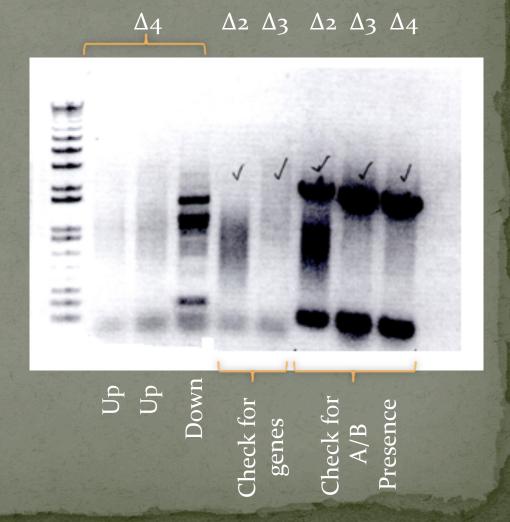
PCR Check

Check for A/B Up & Down

 Δ_2 Δ_3 Δ_4



Up Up Up



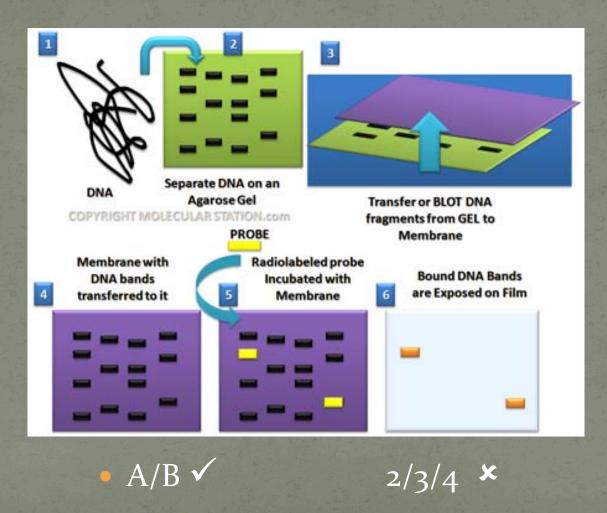
Methods: Analysis of deletion strains

PCR and Transformation 1 and 1 a Transformant ChIP Analysis Grow Tissues RNA Sequencing Southern Western blotting blotting

Phenotype Check

PCP/Check

Southern blotting

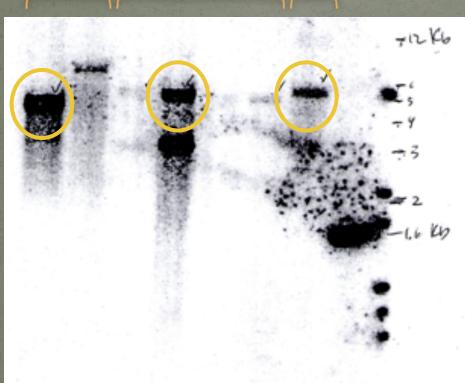


http://enfo.agt.bme.hu/drupal/en/node/8926

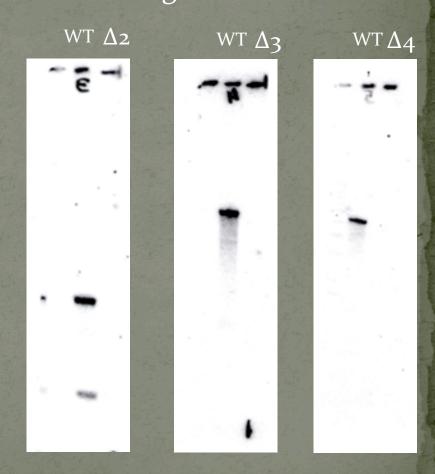
Southern blots

Check for A/B:





Check for genes:



→ Mutants possess A/B

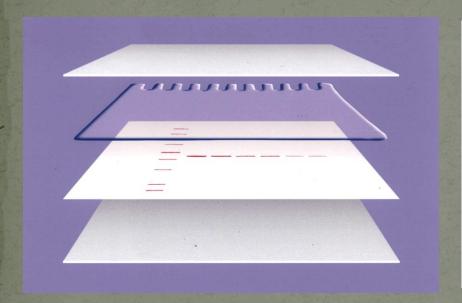
→ Mutants lack KO genes

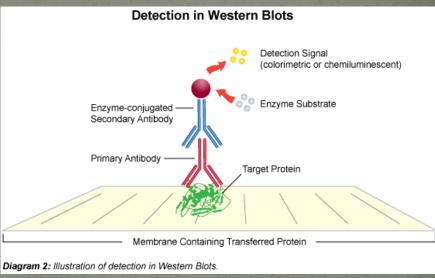
Methods: Analysis of deletion strains

Check

PCR and Transformation 1 and 1 a Transformant Phenotype ChIP Analysis Grow Tissues PCR Check RNA Sequencing Southern Western blotting blotting

Western blotting





- Deletion mutants are predicted to have no H3K27me3 markers
 - Exceptions may be present in $\Delta 2$

http://news.thomasnet.com/fullstory/Membrane-Sandwiches-make-western-blotting-easier-13151 http://www.leinco.com/general_wb

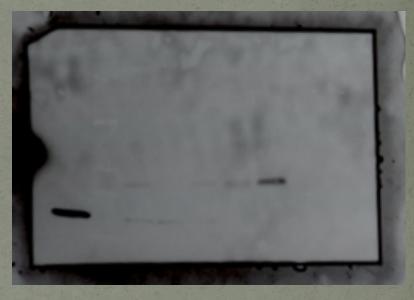
Western blotting

WΤ*
WT
Δ2
Δ3
Δ4



Before water wash

WT WT WT Δ2 Δ3 Δ4



After water wash

- • Δ 3 and Δ 4 do not have H3K27me3 marker
 - • Δ 2 shows weak signal for the marker

Next...

- Investigate by ChIP technique
- Analyze using RNA sequencing method
- Perform gene tagging

Applications

- Learn how PRC2 operates as a protein complex
- Understand importance of each protein subset
- Recognize functions of PRC2 in cell development

Acknowledgments

Dr. Michael Freitag

HHMI

• Dr. Kevin Ahern

URISC

Lanelle Connolly

CURE

Freitag Lab

Oregon State University