

Genetic Mapping and Characterization of the Pale Yellow Locus Responsible for Accelerated Senescence in Dark Tobacco

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Outline

- Background and Application of Pale Yellow (PY) trait
 - Reduced Curing Time
 - Improved Quality
- Mapping of the PY locus
- Genes identified within the Quantitative Trait Locus (QTL) region



Pale Yellow Trait

Found in
accession
TI1372

Accelerate
chlorophyll
breakdown

Trait noticeable
usually by
flowering and
after topping

Traditional
breeding using
phenotypic
selection*

Introduced into
several dark
lines

* Detached Leaf Ethephon Treatment Adapted from the LC Protocol
<https://www.uky.edu/Ag/Tobacco/Pdf/LC-Protocol.pdf>

Chaplin James F.; 1969; *Inheritance and Possible Use of Pale Yellow Character in Tobacco 1*; Crop Science Society of America; 9 (2) p. 169

Application – Reduce Curing Time

Guidelines for Fire Curing Dark Tobacco*



Yellowing

- Ventilation as needed
- None or low heat
- Temp. not to exceed 100°F (37°C)
- 5 – 8 days



Color Setting

- Little or no ventilation
- Temp 100°F-115°F (37°C - 46°C)
- 7 – 14 days



Stem Drying

- Full ventilation
- Temp not to exceed 130°F (54°C)
- 4 – 8 days



Finishing

- No ventilation
- Temp not to exceed 120°F (48°C)
- 10 – 14 days

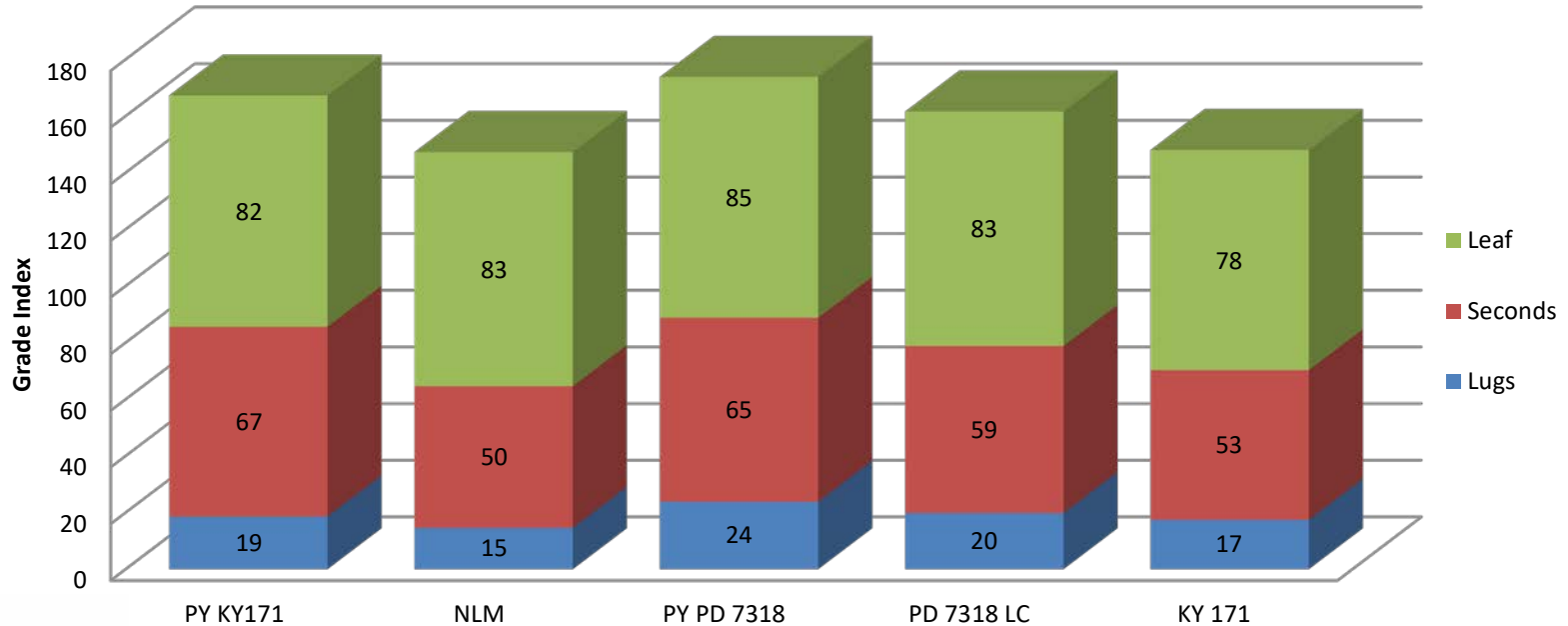


Lusso M., Hayes A., Lion K., Davis G., Hart F., Morris J., 2014; *Methods of Reducing Tobacco-Specific Nitrosamines (TSNAs) And/Or Improving Leaf Quality in Tobacco*; US 2014/0076339 A1

*Bailey, A., 2006. Harvesting, Curing, and Preparing Dark Fire-Cured Tobacco for Market. University of Kentucky – College of Agriculture, AGR152.

Application – Improved Quality

Grade Index (GI) Average of Two Locations in 2009: Blackstone, VA and Princeton, KY



Mapping of Pale Yellow Locus

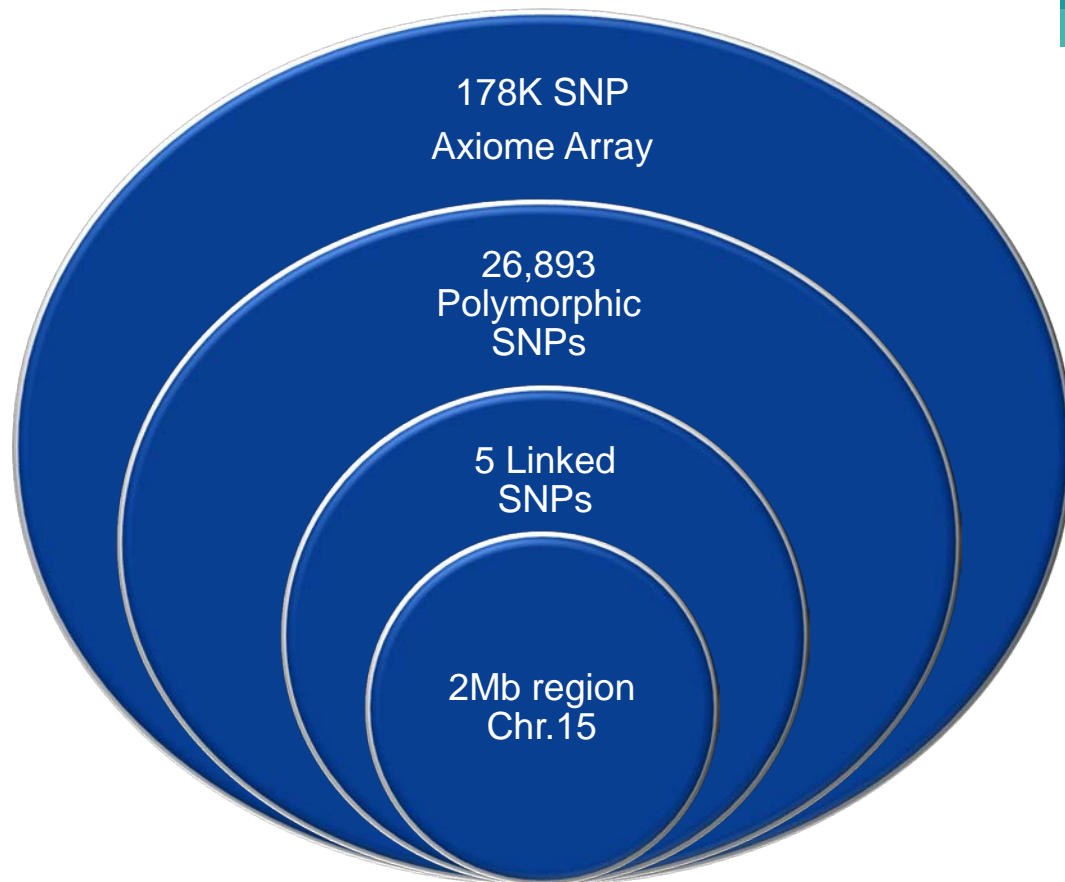
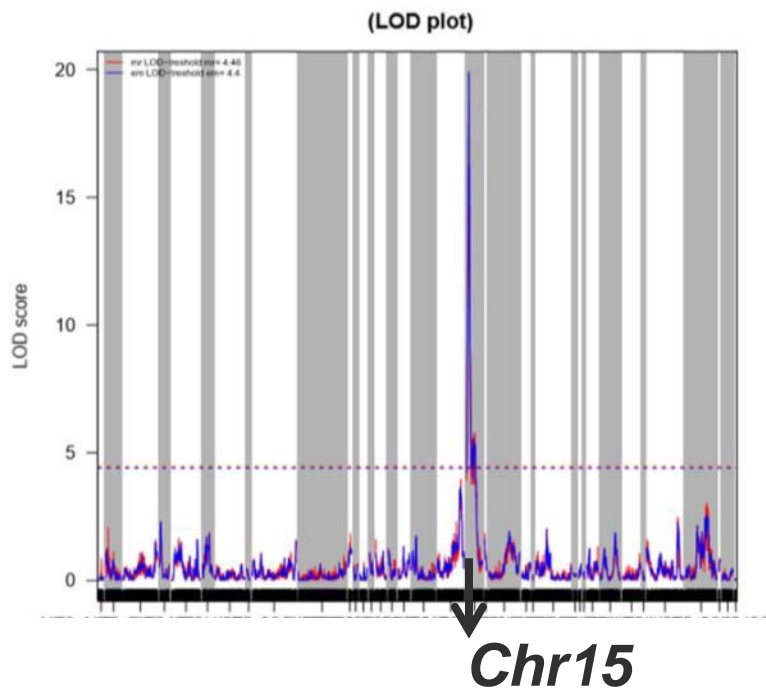
A 192 individual F2 population (NLM x TI1372)x

All individuals were scored on phenotype by

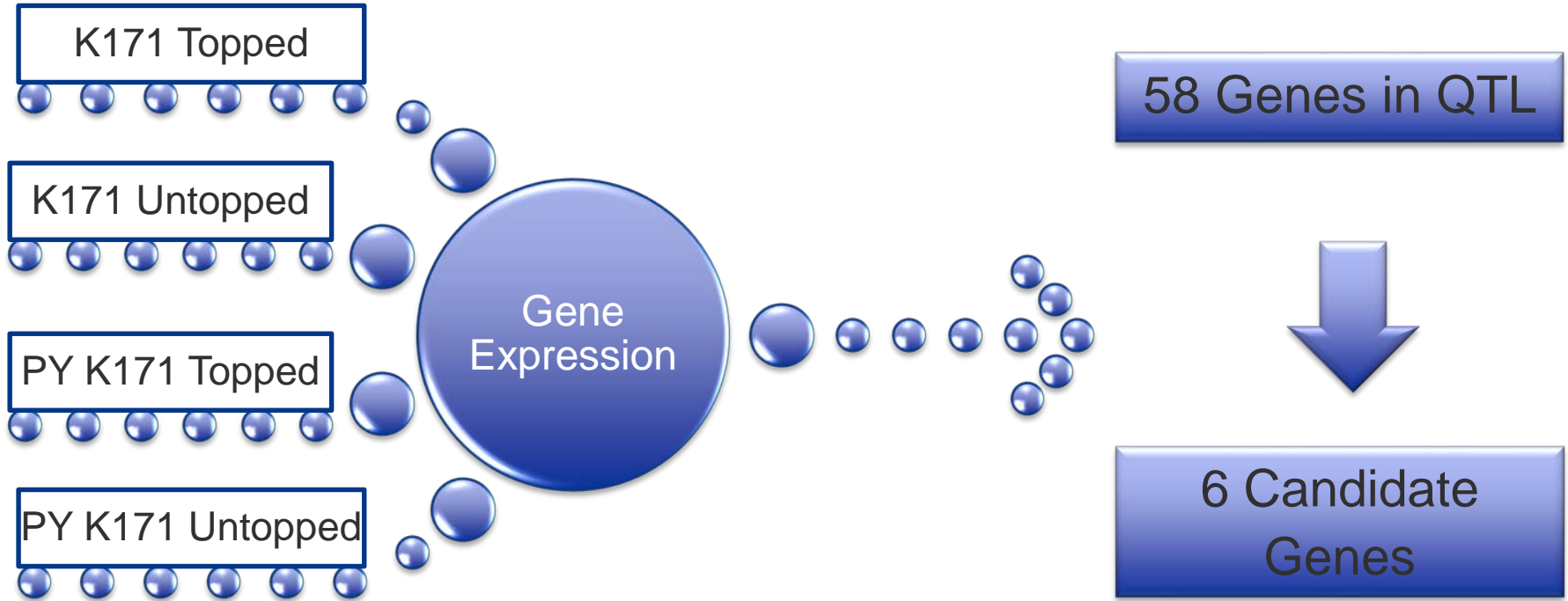
- Visual observation at 4 weeks post topping
- Ethephon screening of leaves at flowering leaves- 5 readings post treatment

93 F2 individuals and parents genotyped on the ~178K SNP tobacco Axiome array

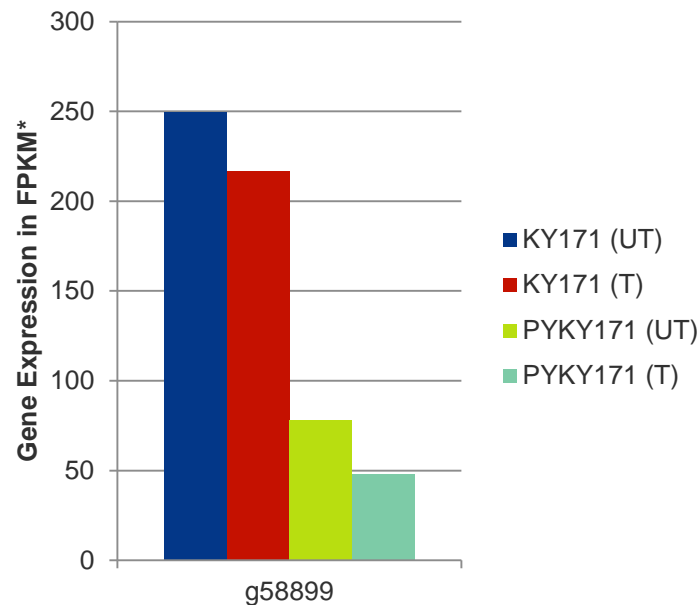
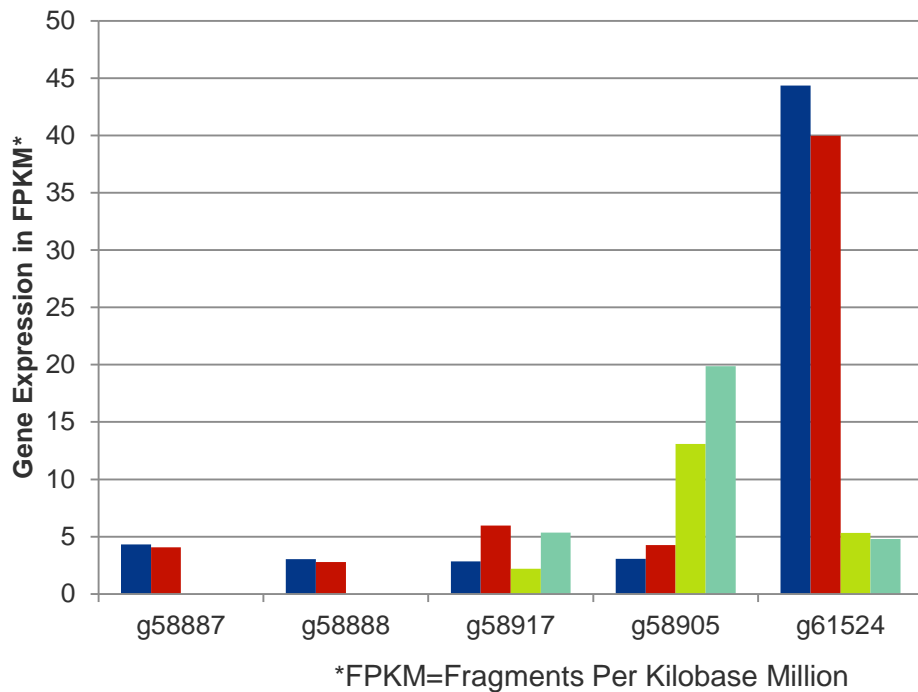
QTL Analysis



Identification of Gene Candidates



Expression Profile of Candidate Genes



Conclusions and Future Directions

Markers

- Five SNP markers for PY trait breeding
- Six functional gene candidates
- Further screening for functional gene marker

Gene Validation

- Allele sequencing
- Gene knockdown / overexpression
- Loss of function mutants from EMS populations



Thank You

QUESTIONS

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