### Genetics of Arabinoxylans

# Characterization of ways to measure arabinoxylans

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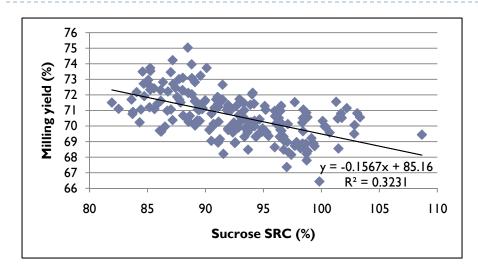
USDA ARS Laboratories Wooster OH and Ohio State University Research funded by USDA-ARS and Kraft Foods



# Background

- Water soluble arabinoxylans (WE-AX) and arabinogalactans (AG) are collectively non-starch polysaccharides that contribute to the sucrose SRC values and water absorption of flour
- AX, when complexed, form insoluble hemi-cellulose material that comprise a significant part of the cell walls of the endosperm and bran
- The quantities of WE-AX and AG in flour are the result of genetics, environment, and flour mill

#### Sucrose SRC and Milling Yield are Intertwined in Soft Wheat



78 76 Milling yield (%) 74 72 70 68 66 64 = -0.0733x + 77.487 62  $R^2 = 0.0723$ 75 85 95 105 115 125 Sucrose SRC (%)

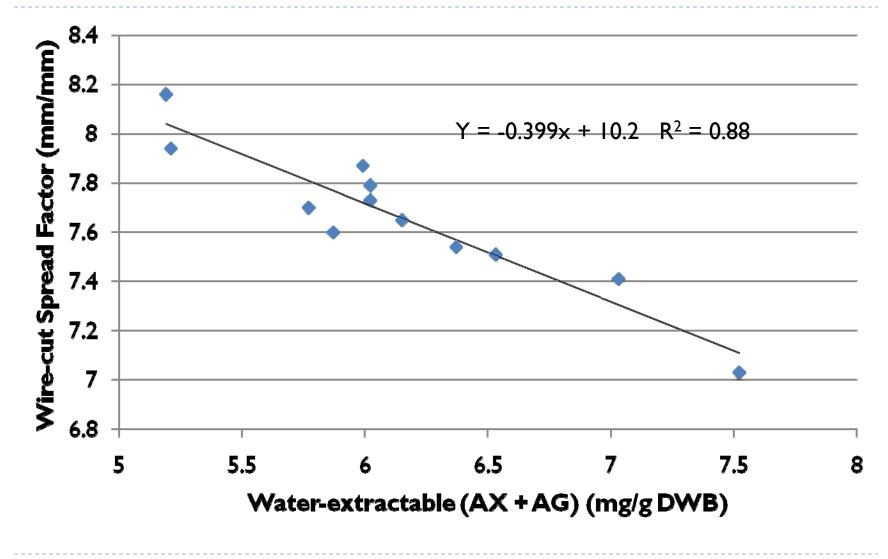
182 soft wheat cultivars Association Mapping Study Genetic means for each wheat grown across 7 environments

#### Genetic correlations are strong

182 soft wheat cultivars Correlation of individual observation each cultivar within each of the 7 environments.

<u>Environmental and phenotypic</u> <u>correlations are less strong</u>

#### Effect of Water-Extractable NSPs on White Flour Wire-Cut Cookies



Guttieri et al. 2008. J.Agric Food Chem. 10929-10932

# Background – Characterizing one level of the interaction

- The quantities WE-AX and AG in flour are the result of genetics, environment, and flour mill
  - Sucrose SRC is a catch-all for a wide variety of compounds
  - Concentration of WE-AX and AG is controlled by genes that synthesize, remodel, and cross-link.
  - As background to studying the gene (next 5 year project plan): characterization of milling system (prior 5 year project plan)



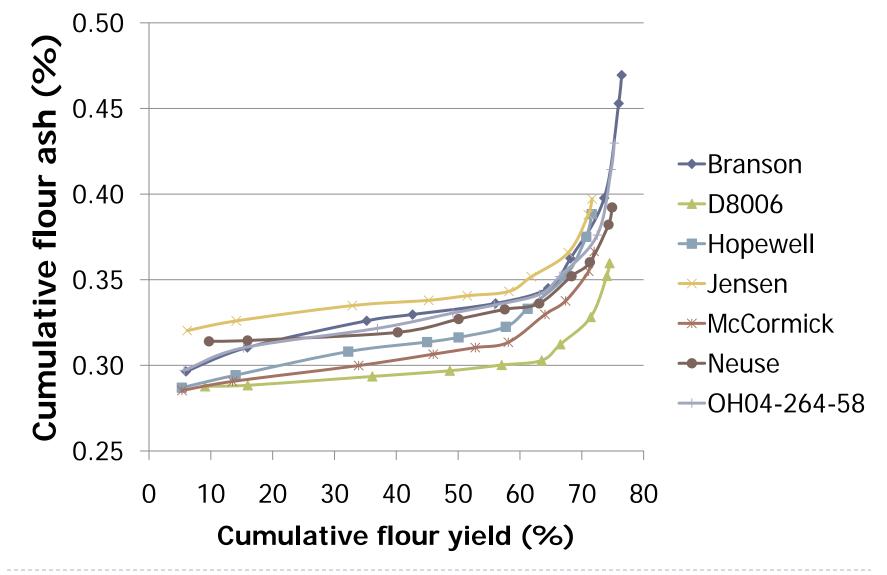
Reduction and bran streams from Miag mill

# Methods (See poster for details)

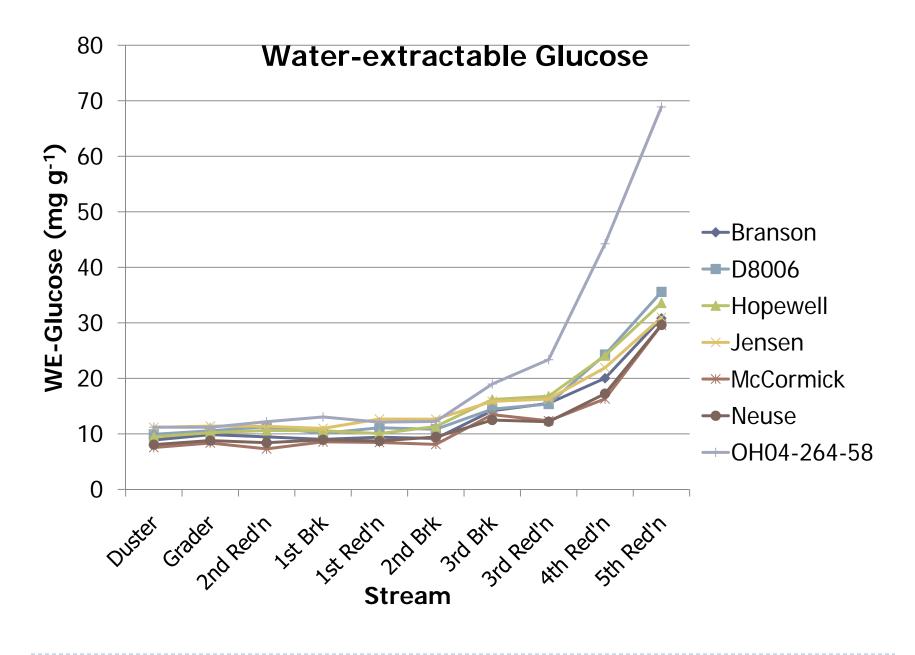
- Seven soft wheat cultivars grown in Wooster, 2009
- Milled on Miag Multomat flour mill
  - I0 streams: 3 breaks, 5 mids, and two resifting streams
  - Mill streams captured separately as in an mill stream
  - Flour ash was analyzed by AACC 08-01
- Water extractable non-starch polysaccharide analysis
  - Aqueous phases from 1 g water SRC of millstreams
  - Hydrolyzed in 2 N trifluoroacetic acid for 1 h at 105 C
  - Derivatized to alditol acetates
  - Separated and quantified by gas chromotography (Guttieri et al. 2008).



#### **Cumulative Flour Ash**



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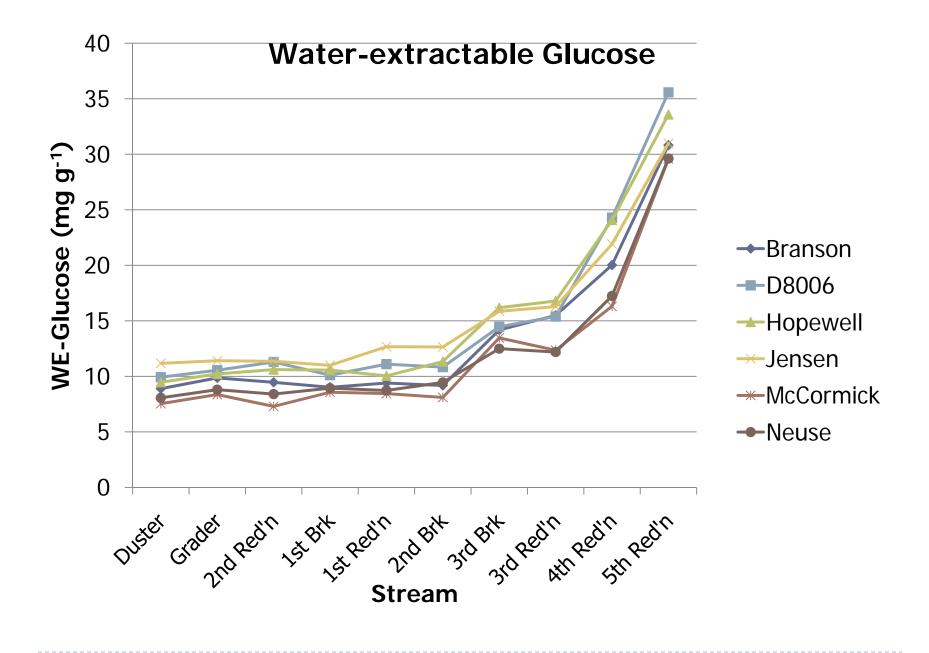


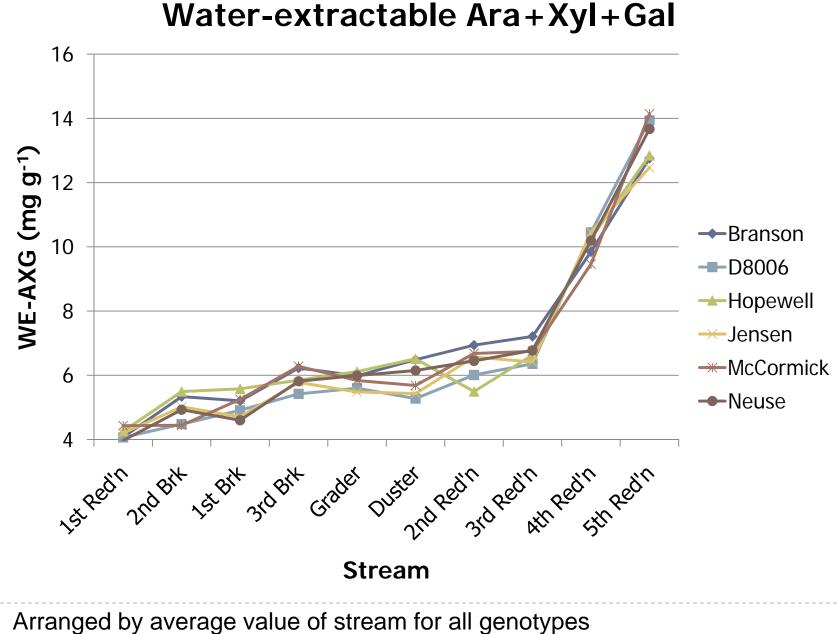
Arranged by average value of stream for all genotypes

# Damaged starch as measured by WEglucose in GC analysis

- Cultivars have interactions with mill streams for the amount of damaged starch that is created in the milling process.
  - OH04-264-58 had the greatest interaction

- For clarity it will be dropped from most of the subsequent slides
- Damaged starch in early roll streams may be independent of later streams.
  - Most damaged starch is derived from 3<sup>rd</sup> break and 3<sup>rd</sup> to 5<sup>th</sup> reduction
  - Greatest sample differences are observed in 3<sup>rd</sup> to 5<sup>th</sup> reduction streams

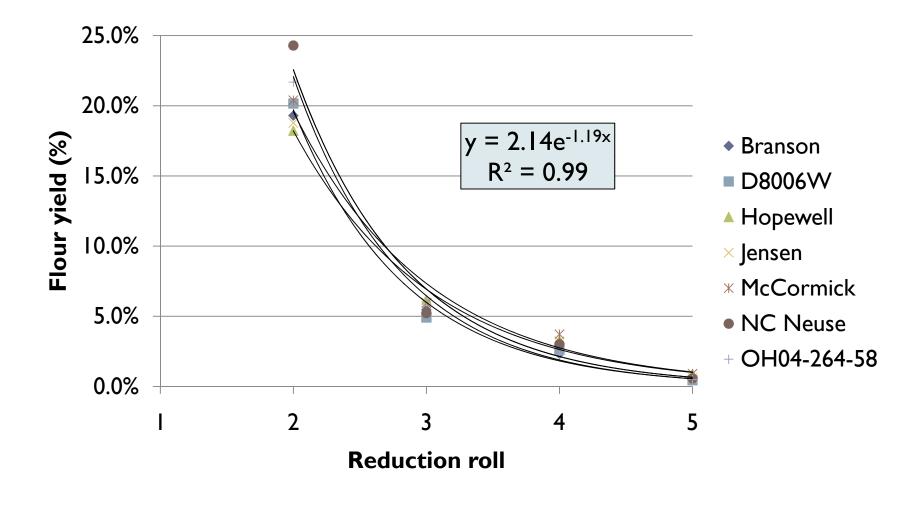




Where do the WE-AX and AG come from in the milling process?

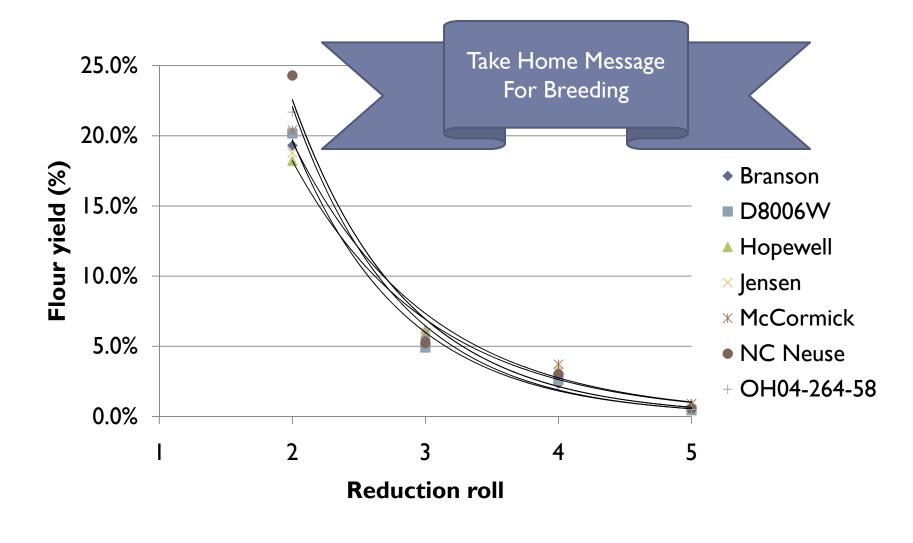
- Total concentration of combined ara, xyl, and galactose is similar among the six cultivars for each stream
- Concentration is greatest in 3<sup>rd</sup> to 5<sup>th</sup> reductions
- Relative differences in total arabinoxylans and arabinogalactans in flour are therefore due to primarily to quantity of flour recovered in different streams
- Structure and composition of the non-starch polysaccharide fraction affect greatly its water activity and impact of soft wheat quality

#### Flour Yield of Reduction Streams



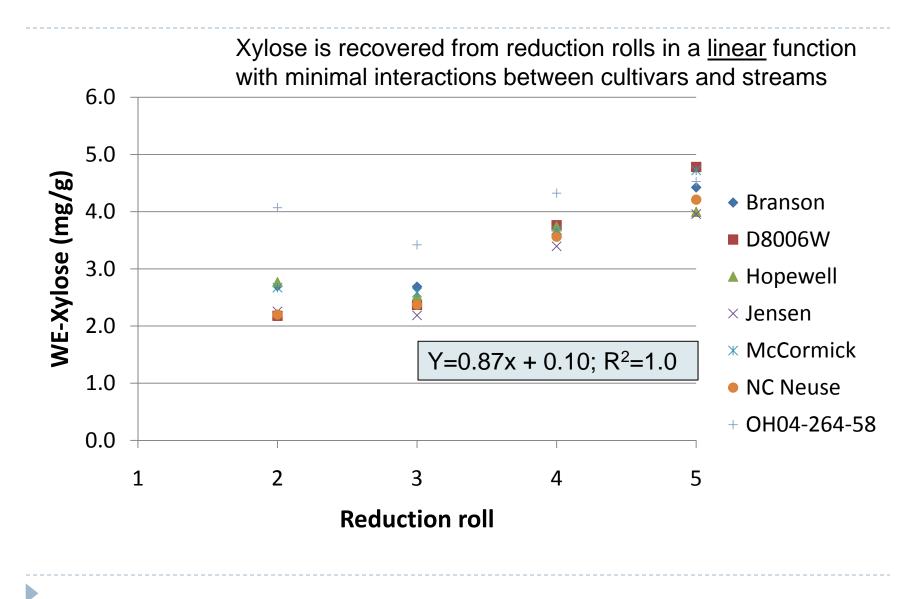
Exponential decay function Minimal differences in flour yield for 3<sup>rd</sup> to 5<sup>th</sup> Red'n

### Flour Yield of Reduction Streams

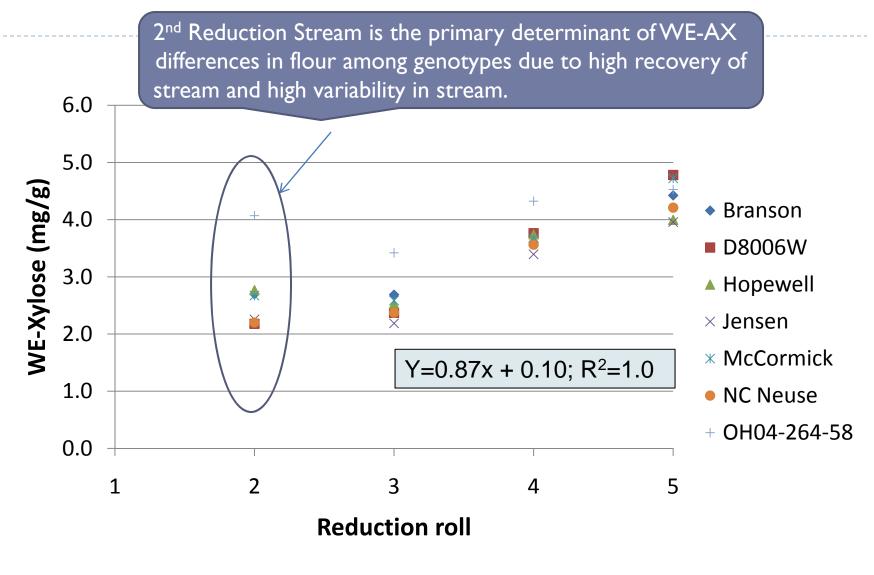


Exponential decay function Minimal differences in flour yield for 3<sup>rd</sup> to 5<sup>th</sup> Red'n

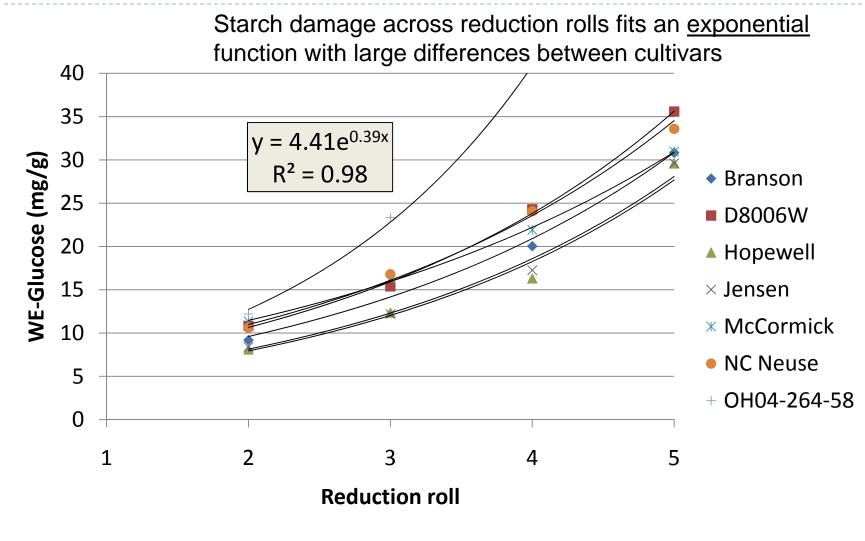
#### WE-Xylose in Reduction Streams



# WE-Xylose in Reduction Streams



#### Starch Damage in Reduction Streams



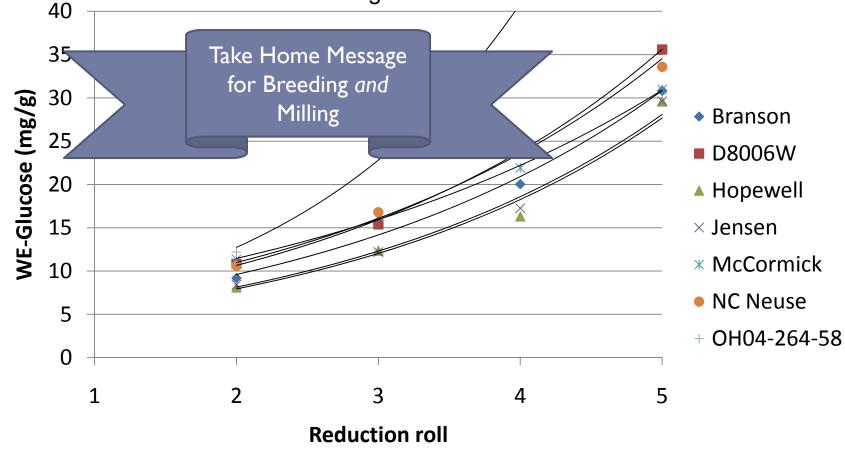
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The exponential function is used to model phenomena when a <u>constant change</u> in the independent variable gives the same <u>proportional change</u> (increase or decrease) in the dependent variable.

www.wikipedia.org

### Starch Damage in Reduction Streams

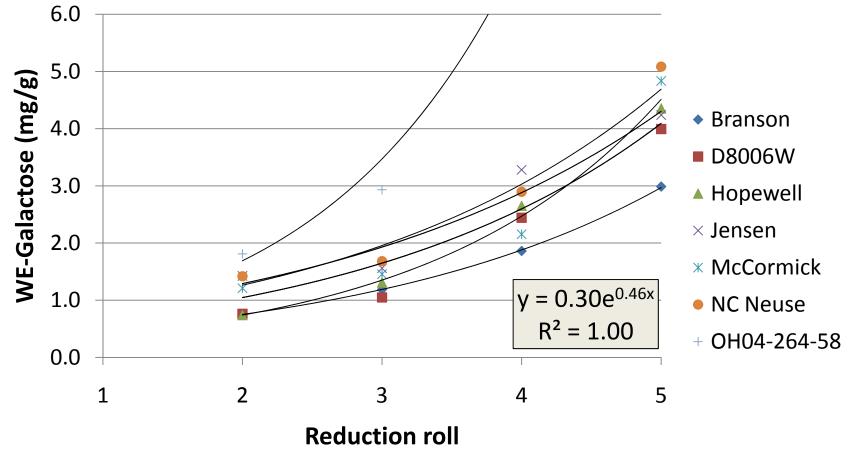
Starch damage across reduction rolls fits an <u>exponential</u> function with large differences between cultivars



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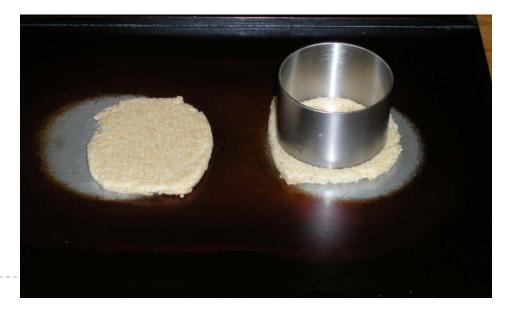
### Galactans in Reduction Streams

Galactans have the greatest variation among cultivars, and the largest interactions between cultivar and rolls. Also an <u>exponential</u> function.



# Galactans in milling and flour

- Galactans are typically present in the flour in the form of arabinogalactans (AG) and are often bound to short peptides cleaved from the Grain Softness Protein
  - AG have limited correlations to sucrose SRC values
  - AG disproportionately reduce soft wheat flour quality for high sugar baked products like cookies



Wire-cut cookies made with whole wheat flour

# Galactans in milling and flour

- AG peptides in other tissues and other plants affect cell wall rigidity
  - Do they control milling behavior?
- Subsequent genetic studies
  - Genes affecting AG synthesis and location
  - Differences in AG content in reduction flour may be diagnostic

Arabinogalactan peptides visualized with Yariv stain in wheat endosperm



# Conclusions for Milling

- The amount of flour recovered in later reduction rolls is largely a mechanical function rather than a grain function
  - Quad micro mills predict long-flow mills well because of this relationship
- Although flour quantity is determined by the mill in later reduction rolls, the damaged starch caused by the recovery and arabinogalactans added to flour varies greatly among cultivars
- Quality of flour recovered decays much faster than the amount of flour recovered in later reduction rolls due to compound action of rolls on particles resulting in exponential damage



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