

What the Dickens has Oat Genomics Done for <u>You</u> Lately ?

The Groats of Christmas <u>Past</u>, <u>Present</u>, and <u>Yet-to-Come</u>

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First, please appreciate the Oat Genome – in context:

12.7 Billion bp / 2 x 21 chromosomes = 6 metres of DNA

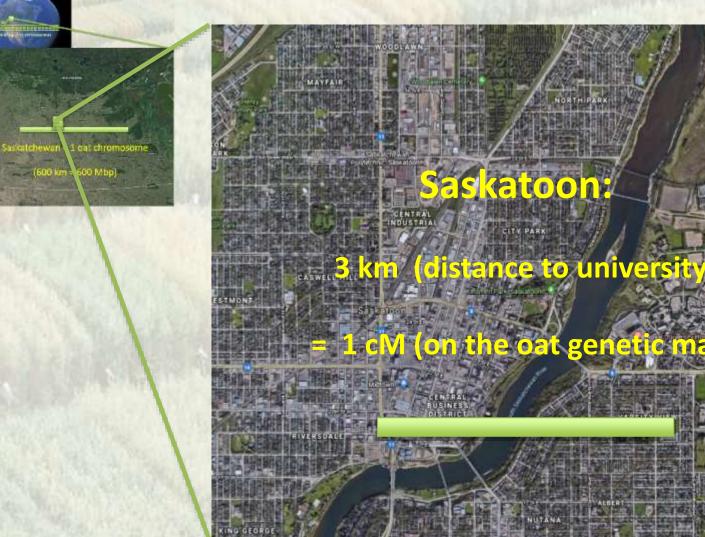
First, please appreciate the Oat Genome – in context:



Stretched to 12,742 km (1bp = 1mm)

12.7 Billion bp / 2 x 21 chromosomes = 6 metres of DNA





= 1 cM (on the oat genetic ma 0 MANAGEMENT



1. The oat genome is huge and complex

.....

2. You need to know where to look, and what you are looking for

> Statisticon Mutations Asso

1 person (2m) 1 sene (2000 bp)

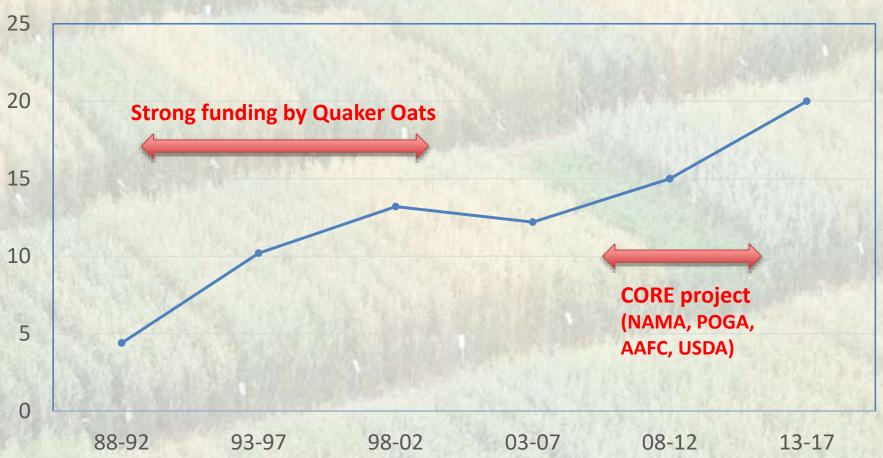
Saskatch

The "groats of Christmas Past"

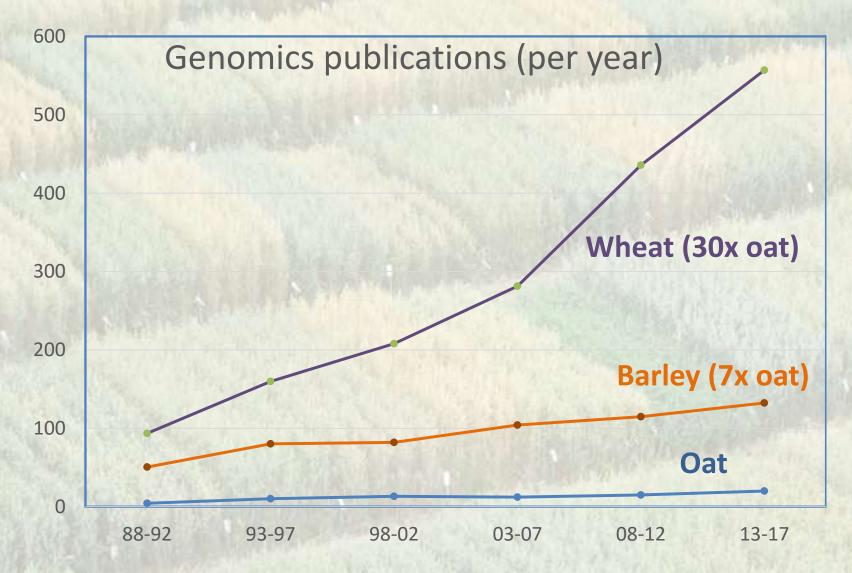


30 years of oat genomics

Oat Genomics publications (world-wide, per year)



Reality check.....



What have we accomplished ?

- Essential genomics Infrastructure
 - Integrated genetic map (despite complex genome)
 - Molecular markers (as good as other crops)
 - QTL studies (map locations of genes)
- Marker assisted selection for key rust genes
- Understanding and broadening of germplasm
- Complete genome sequences in progress
 - Wild relatives (smaller genomes)
 - Cultivated oat (US nearly complete + Swedish group...)

The 'past' should not be 'passed'

Charlene Wight, AAFC



Oat Newsletter (OatNews.Org)



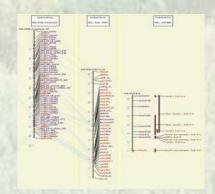
Pedigrees of Oat (Pool.aowc.ca)

1	2	3
	Putoam.	Purdue 3714A4-7-5
Brave		Clinton
	unnamed_2618	unnamed_6710
		Andrew
	Ivier	Clintland 60

T3 Oat (triticeaetoolbox.org/oat) Unpublished data sets

More Select - Analyze - Download - Mouse e More Select - Analyze - Download - Mouse e Marcial Communication Select - Analyze - Download - Mouse e Marcial Communication Select - Communication Marcial Communication Select - Select

GrainGenes (wheat.pw.usda.gov) Published data

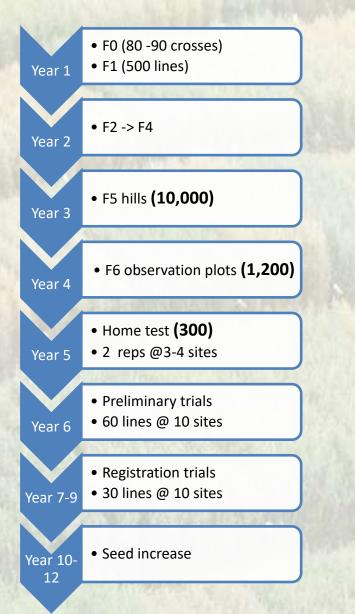


The Groats of Christmas Present



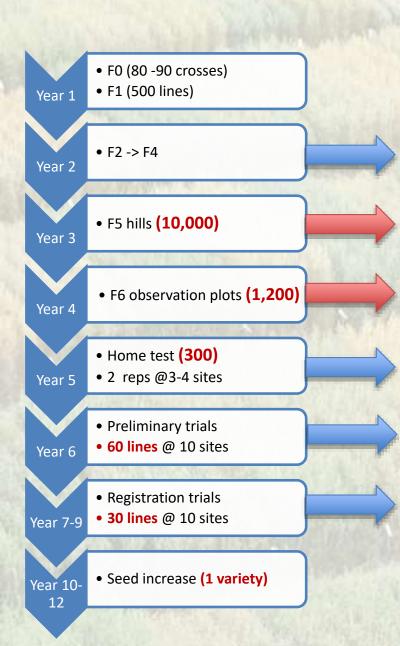
* Focus on genomic selection (largest impact)

Plant breeding takes 10 to 12 years



Weikai Yan – Oat Breeder, AAFC Ottawa



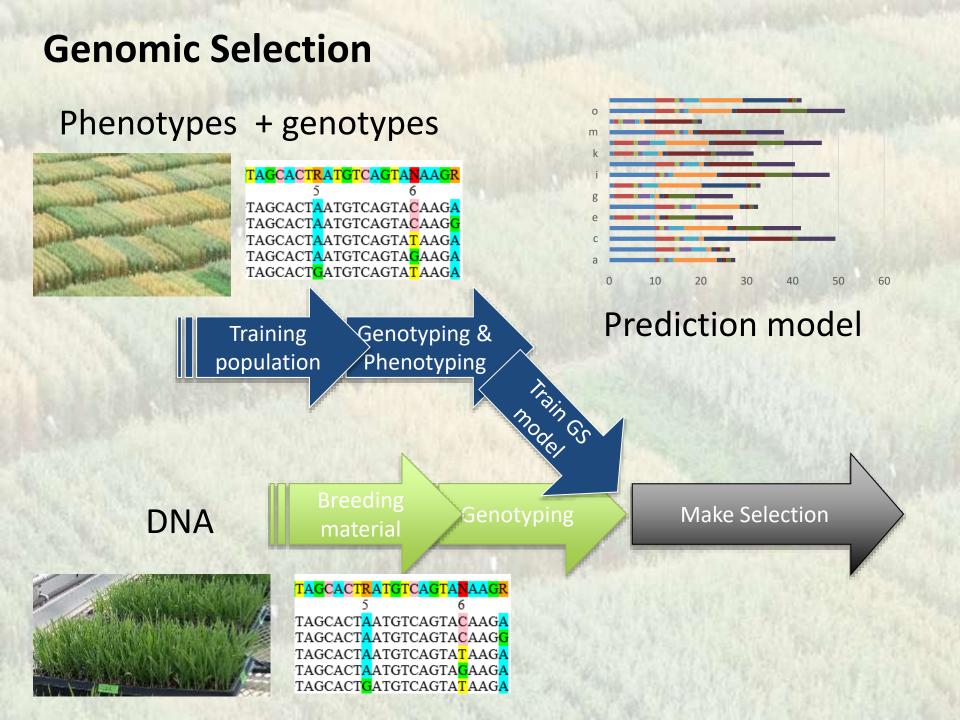


10,000 potential oat varieties.....

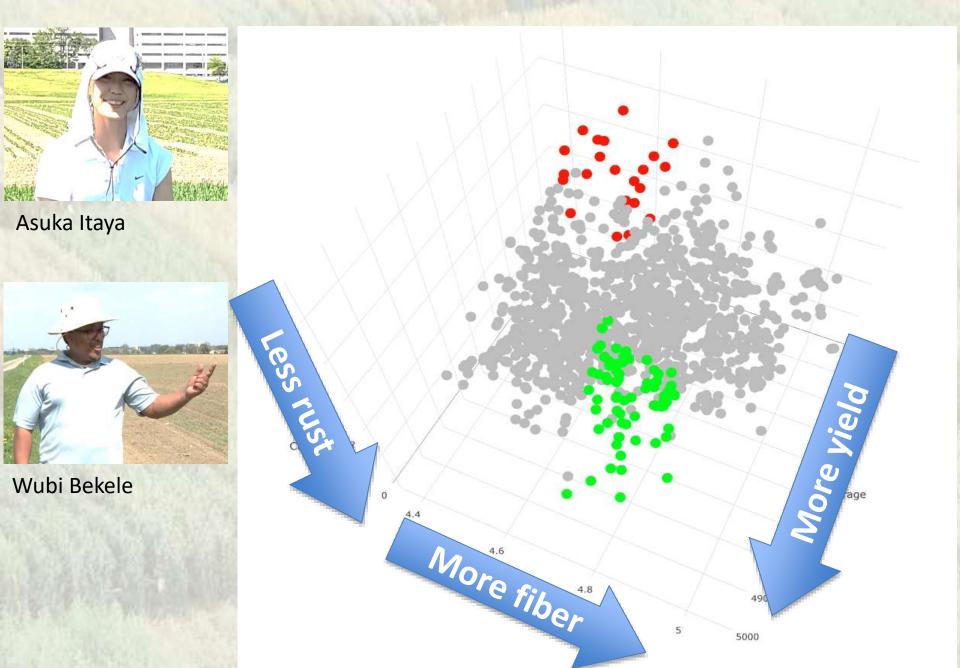


Mostly visual selection

Multi-year Multi-location Large plots

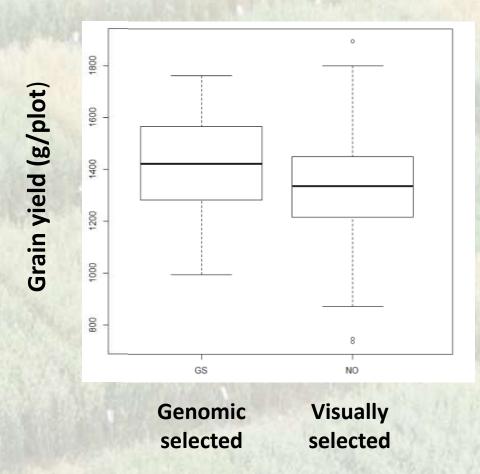


Genomic selection (2018 - 3 traits)



Genomic selection: Validations (2016 \rightarrow 2018)

Lines selected based on DNA show higher yield than those selected visually



The Groats of Christmas Yet to Come...



Goals

Genomic selection:

- Faster, cheaper, more automated

- E.g. next-year's GS will cost < \$15 per sample
- Prediction is better through more training environments (and experience)
- GS is cost-effective and indispensable

Goals (continued)

- Oat is fully sequenced (more than once)
- Historical info and model species link to oat genome sequence database
 - What do genes do ? Why are they important ?
- Gene diversity database
 - What versions of these genes exist in oat ?
- Gene editing (for gene validation)
- Genomic selection integrated with high-value target genes...
 - What do I mean by this??

Genomic selection vs target genes

1990 - 2010:

Map genes and select them one-by-one
2010 - 2015:

This is not working very well!

2015-2020:

- Genotype the whole genome
- Statistical models, weighted markers
- Integrates well with breeding

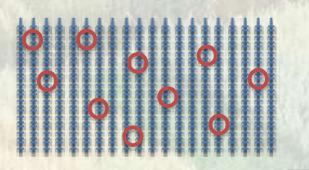
2020-2030?

- In depth biochemistry & physiology
- Identify gene versions with large effects
- Combine with knowledge from genomic selection
- = Integrative genomic breeding

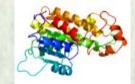


Quantitative Trait Locus (QTL)

- Small part of one trait
- limited accuracy







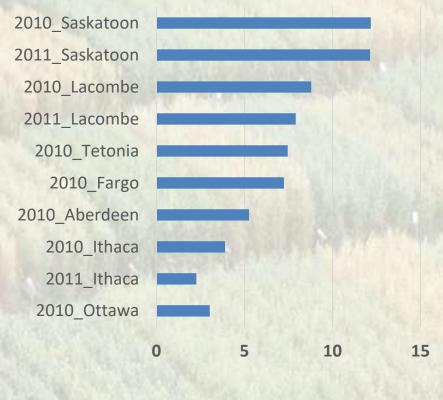
Example: marker "avgbs_201782"

TGCAGAGGAACGGAGGCAGTATTGTACTACATGYACTCCTGTACCTCTCAGCAGTCACCGACAC 1 2 3 4 5 6 TGCAGAGGAACGGAGGCAGTATTGTACTACATGCACTCCTGTACCTCTCAGCAGTCACCGACAC TGCAGAGGAACGGAGGCAGTATTGTACTACATGTACTCCTGTACCTCTCAGCAGTCACCGACAC

Every oat we have tested has either the "C" or "T" version

Effect of "C" vs "T" (averaged over 400 spring oats)

"C" makes oats yield more



% yield increase (C vs T)

"C" makes oats head later



Days to heading (C vs T)

5

avgbs_201782

TGCAGAGGAACGGAGGCAGTATTGTACTACATGYACTCCTGTACCTCTCAGCAGTCACCGACAC 1 2 3 4 5 6 TGCAGAGGAACGGAGGCAGTATTGTACTACATGCACTCCTGTACCTCTCAGCAGTCACCGACAC TGCAGAGGAACGGAGGCAGTATTGTACTACATGTACTCCTGTACCTCTCAGCAGTCACCGACAC

- Avgbs_201782 = linked to a gene that responds to long days.
- I hope our future work will prove this, and characterize the gene
- Then what. Who cares?
 - Often we cross with germplasm containing the "T" version
 - Then we need to re-select the "C" version
 - Genomic selection can accomplish this
 - But if we are deliberate in our selection of this gene, GS can be more efficient in dealing with other genes.

Traits where one gene can make a difference

- Rust (AAFC/USDA)
- Dwarfing (AAFC/UK)
- Flowering time (AAFC)
- B glucan and oil (AAFC/USDA/Cornell)
- Drought (AFFC/Spain)







Value of oat genomics in Canada ???

1% yield increase = \$ 10 M 1% value increase = \$ 10 M 1% export increase = \$ 5 M Etc...

Or... (if we don't keep up):

- Cost of losing oat as a commodity ?
- Cost of losing export share ?
- Cost of losing Canadian-owned varieties ?

What do you think... Let's talk!

Thank You









Canada's Seed Partner





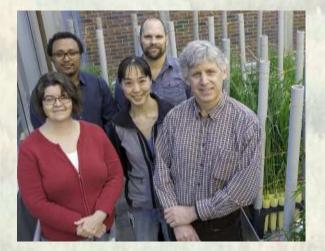
GRAIN MILLERS















Agriculture and Agri-Food Canada