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Canada



Genetic Resources for Pest Management

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Field(s) of Dreams

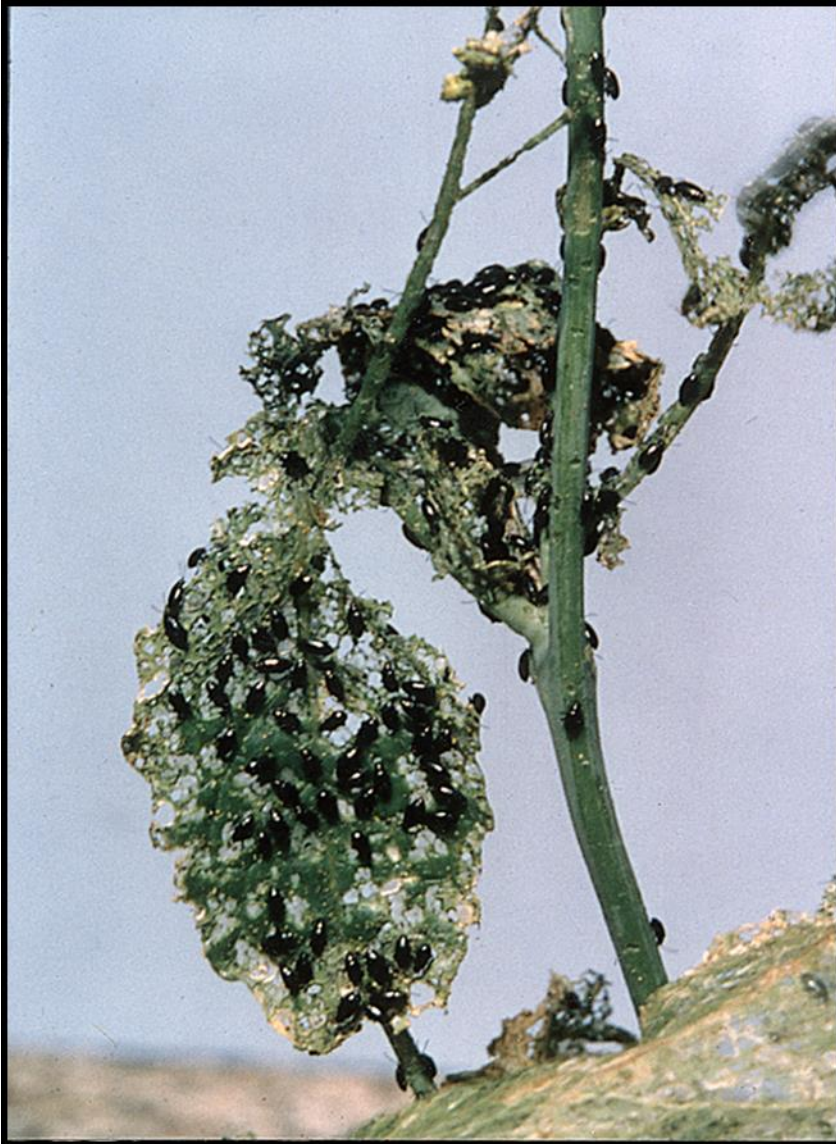


Pl

is



Flea Beetles



Crucifer Flea Beetle

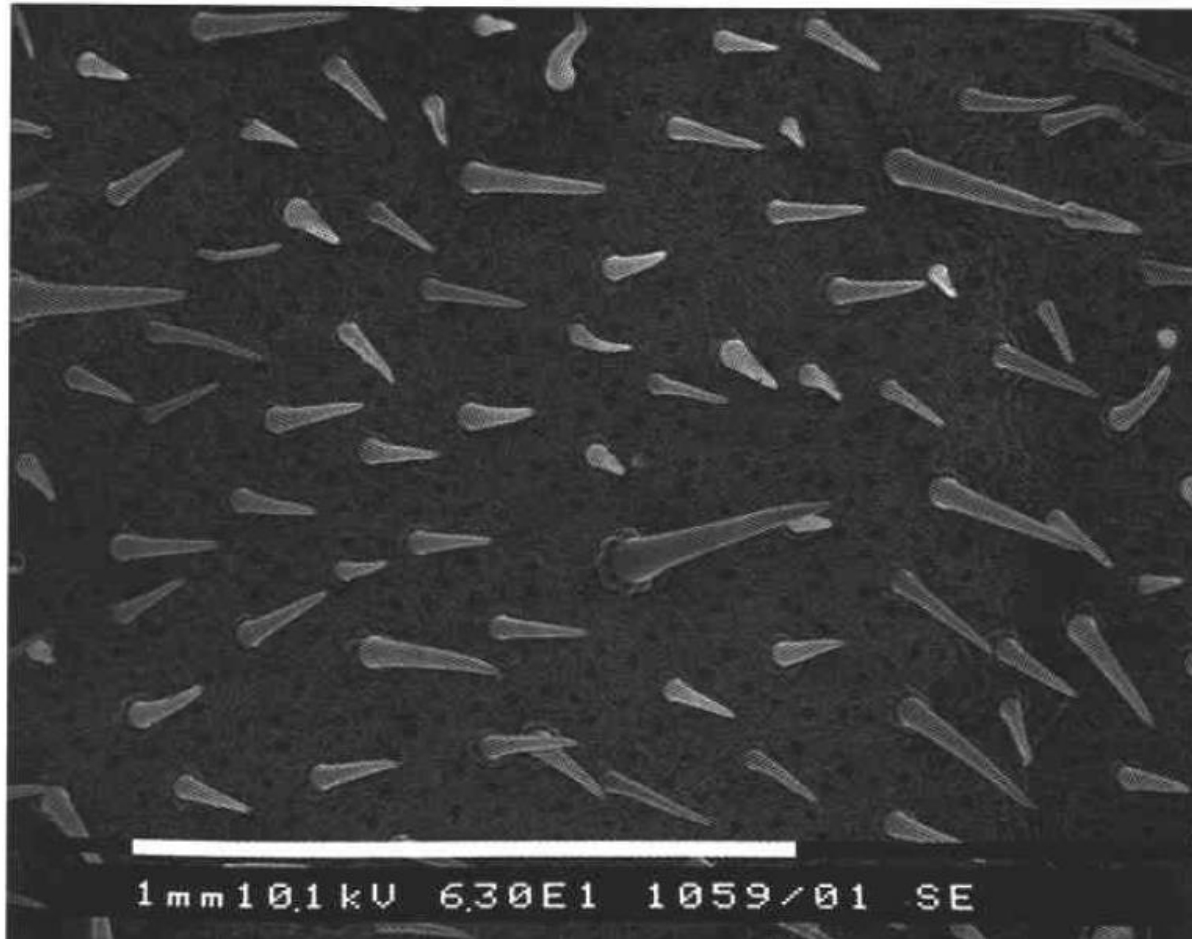


Striped Flea Beetle



Flea Beetles vs Hairs/Trichomes

B. villosa



Palaniswamy P and Bodnaryk R. 1994. A wild **Brassica** from Sicily provides trichome-based resistance against flea beetles, *Phyllotreta cruciferae* (Goeze) (Coleoptera: Chrysomelidae). The Canadian Entomologist 126:1119-1130.

Hairy Canola

- Developed by Margaret Gruber & Julie Soroka (AAFC retired)



Contemporary Canola



Hairy *Brassica napus*

Hairy Canola Disrupts Feeding

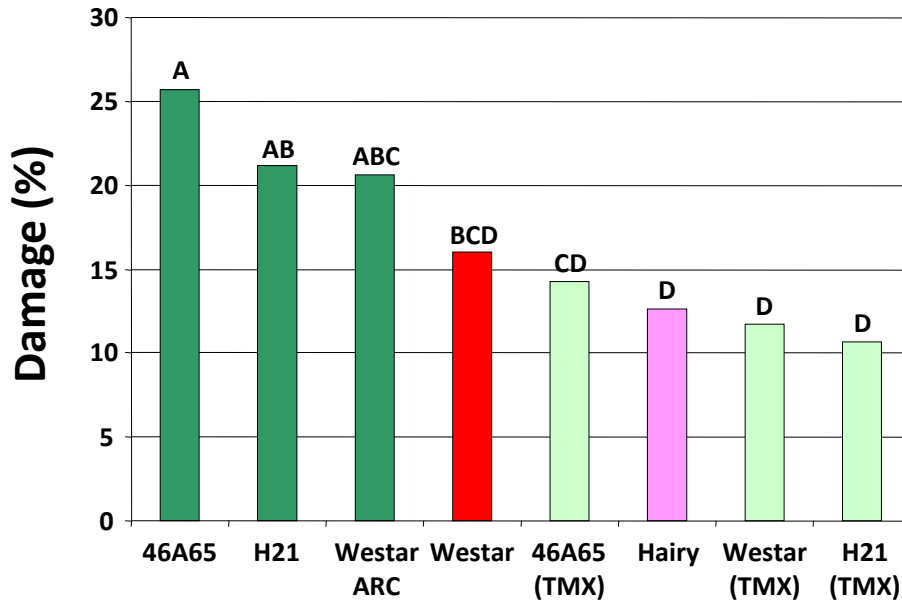


Flea Beetles Avoid Hairy Canola

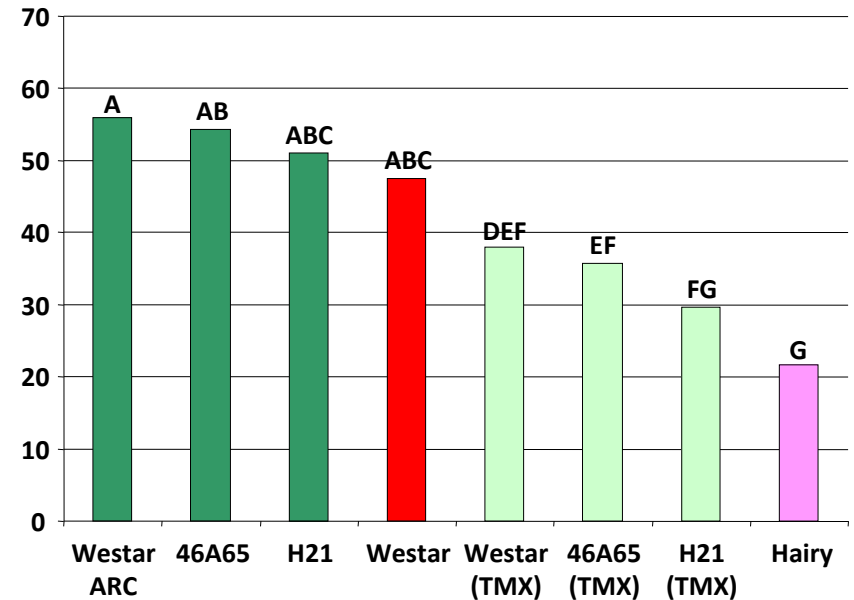


Field Assays

1st & 2nd Leaves – 20 days



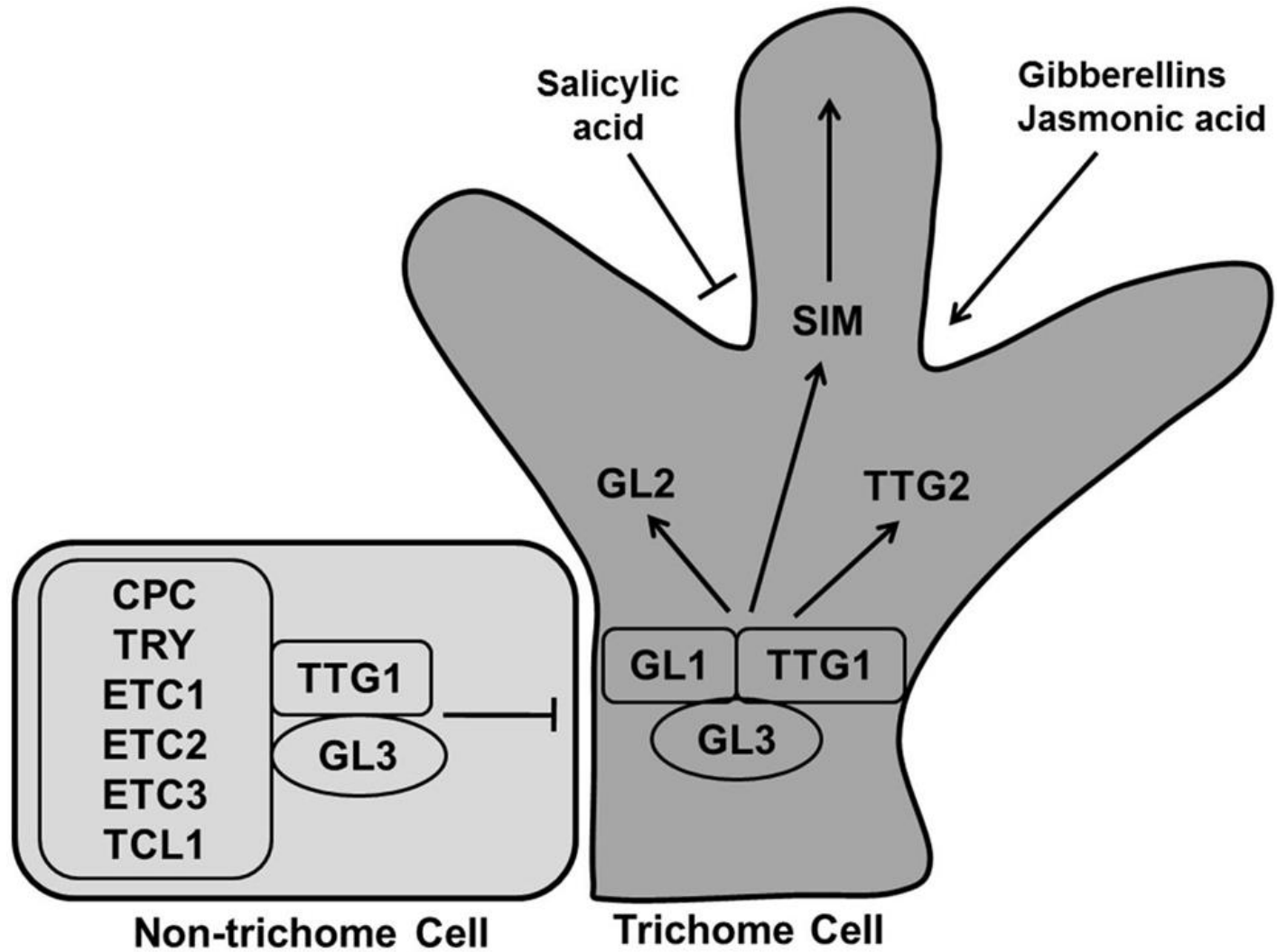
3rd & 4th Leaves – 20 days



TMX = thiamethoxam (Helix)

Soroka J, Holowachuk JM, Gruber MY and Grenkow LF. 2011. **Feeding by Flea Beetles (Coleoptera: Chrysomelidae; *Phyllotreta* spp.) is Decreased on Canola (*Brassica napus*) Seedlings with Increased Trichome Density.** Journal of Economic Entomology, 104:125-136. DOI: 10.1603/EC10151. URL: <http://www.bioone.org/doi/full/10.1603/EC10151>.

How was this achieved?



Replaced GL3 Protein

AtGL3



Arabidopsis thaliana

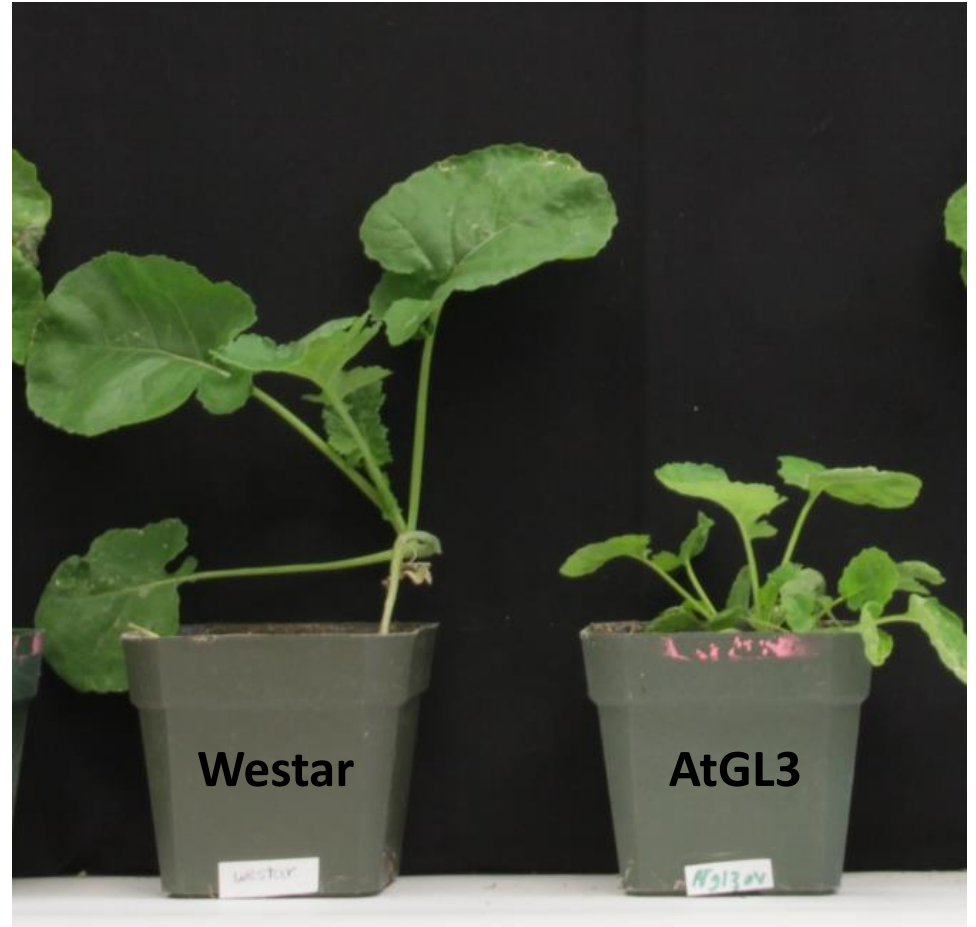


B. napus cv. Westar

What Happened?

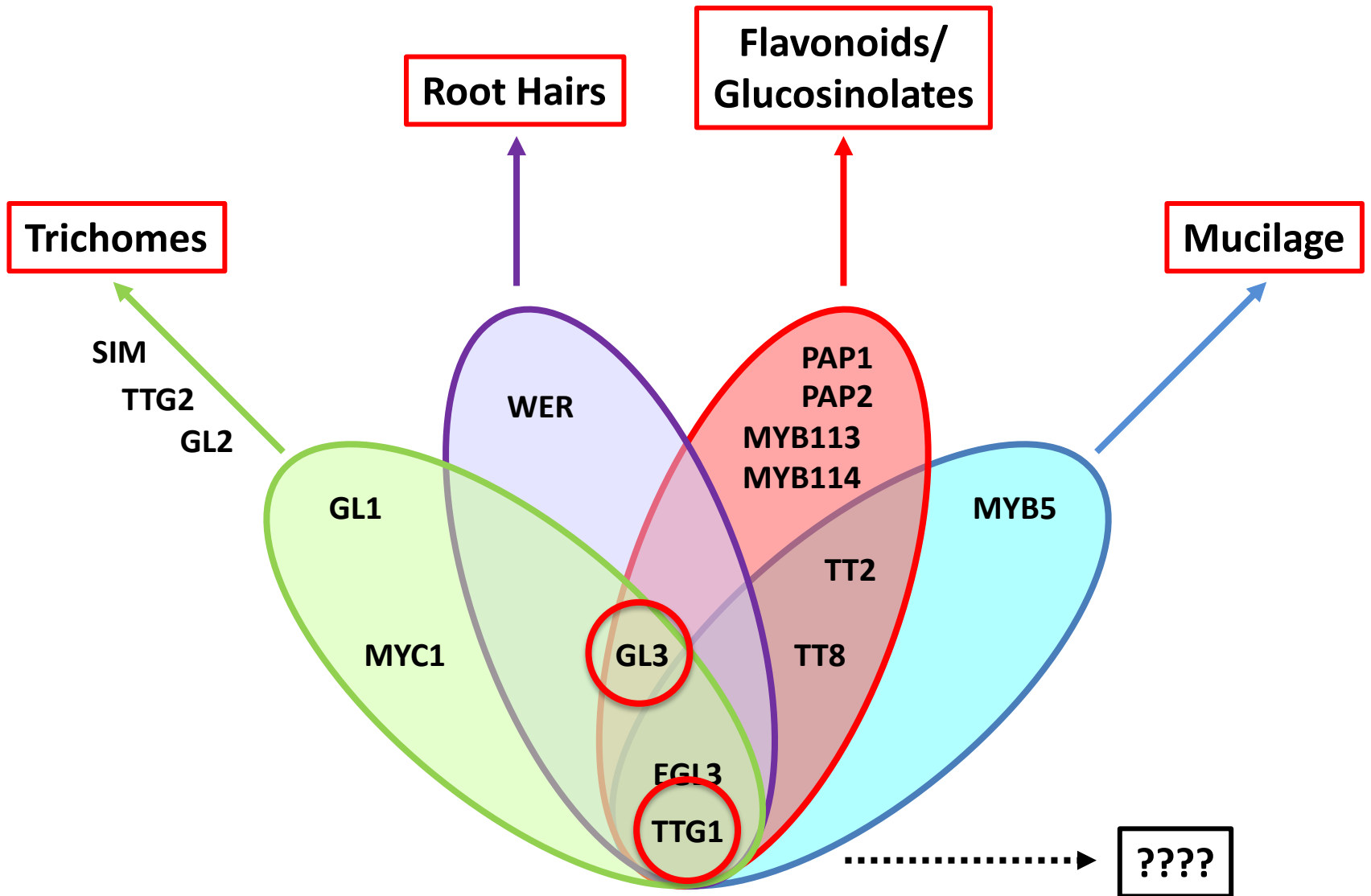


Hair Production



Severe Impact on Growth

Why?



The Fix

- Reduce TTG1 (RNAi interference)



Westar

AtGL3



AtGL3:ttg1
"Hairy Canola"

Why is this not in farmer's fields?

- **Required 2 transgenic events**
 - First event to induce hair production (*AtGL3*)
 - Second event to fix developmental issues (*TTG1*)
- **Patent on *A. thaliana* GL3 (Texas A & M)**
- **Interest from major breeding companies**
 - Registration and regulatory costs for GMO
 - Internal conflict between business units*

What next?

- **Screened 1000 Brassica lines for hairiness**
 - *B. rapa* (A genome - Polish canola – 652 lines)
 - *B. nigra* (B genome – 21 lines)
 - *B. oleraceae* (C genome – vegetables – 26 lines)
 - *B. villosa* (C genome – 1 line)
 - *B. napus* (AC genome – canola – 296 lines)
- Plant Gene Resources for Canada and others

Naturally Hairy Brassicas

“Hairy Canola”



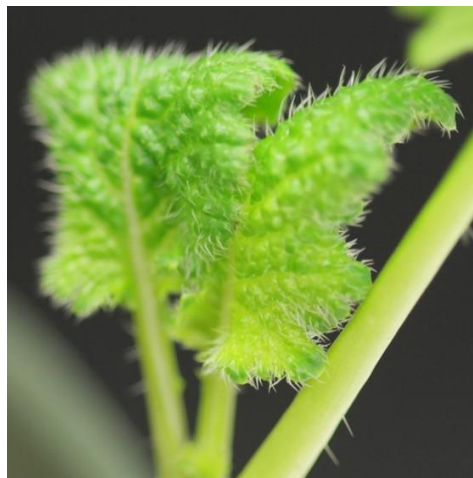
B. napus



B. villosa



B. rapa



Brassica villosa



Nagahabushana NK, Tan Y, Taheri A, Li X, Bjorndahl TC, Nowak J, Wishart DS, Hegedus DD, and Gruber M. 2014. *Brassica villosa*, a system for studying non-glandular trichomes and genes in the Brassicas. *Plant Mol. Biol.* 85:519-539.

Accessing the “Hairy” Trait

1. **GL3 genes from hairy Brassica species**
2. **Map and introgress hairy trait into *B. napus***

Other Brassica *GL3* Genes

Moderately Hairy

B. rapa *GL3*



Hairy *B. rapa* *GL3*



B. villosa *GL3*



Other *GL3* Genes



Westar

***B. rapa* GL3**

***B. villosa* GL3**

- Working on a strategy to fine-tune expression of *GL3* to avoid off-phenotypes

Hairy *B. napus* Accessions



**Naturally
Hairy
*B. napus***

Westar

AtGL3

AtGL3:ttg1

B. napus Crosses

Hairy *B. napus* (DH-Dos2)

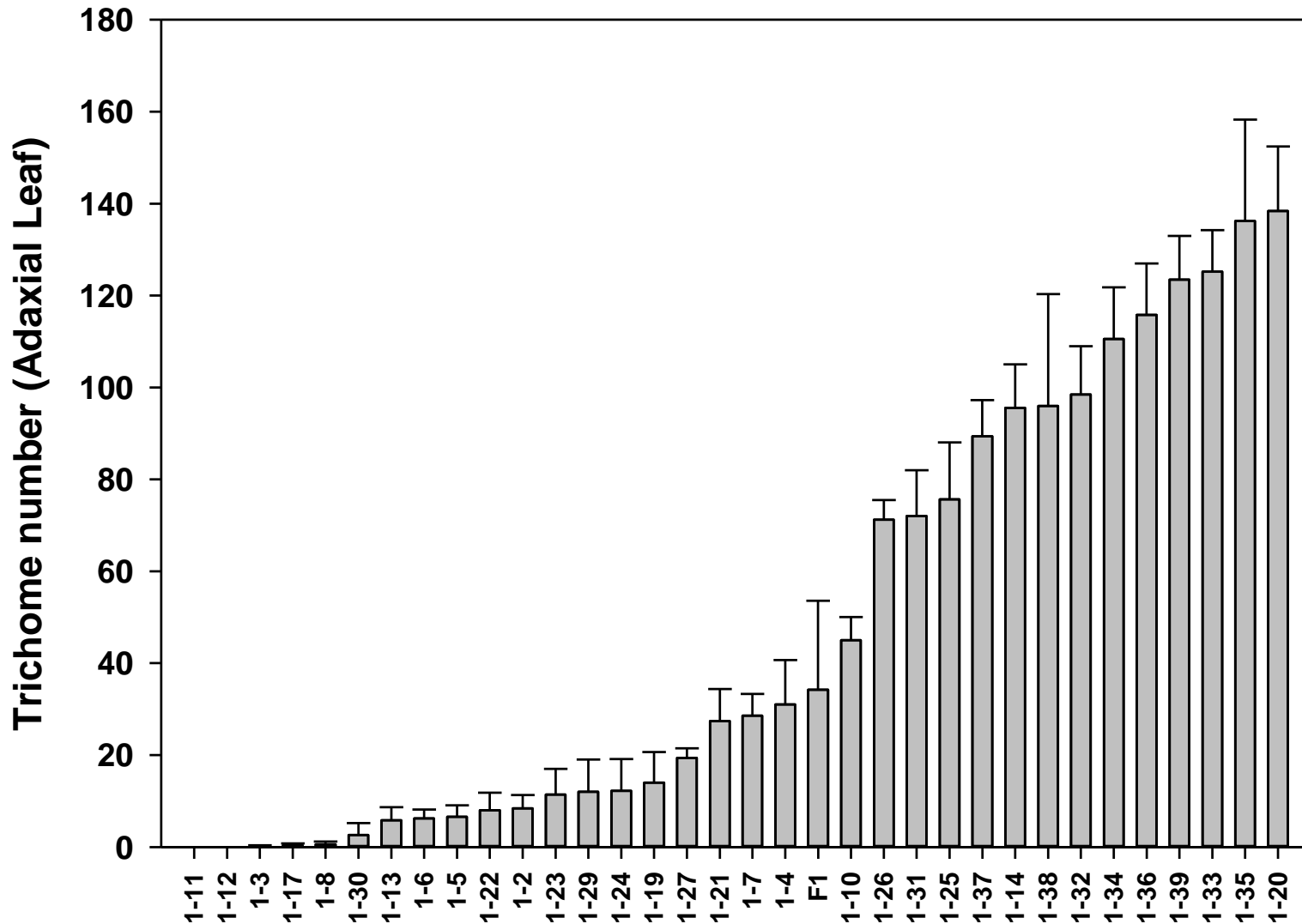


Glabrous *B. napus* (NAM-0)

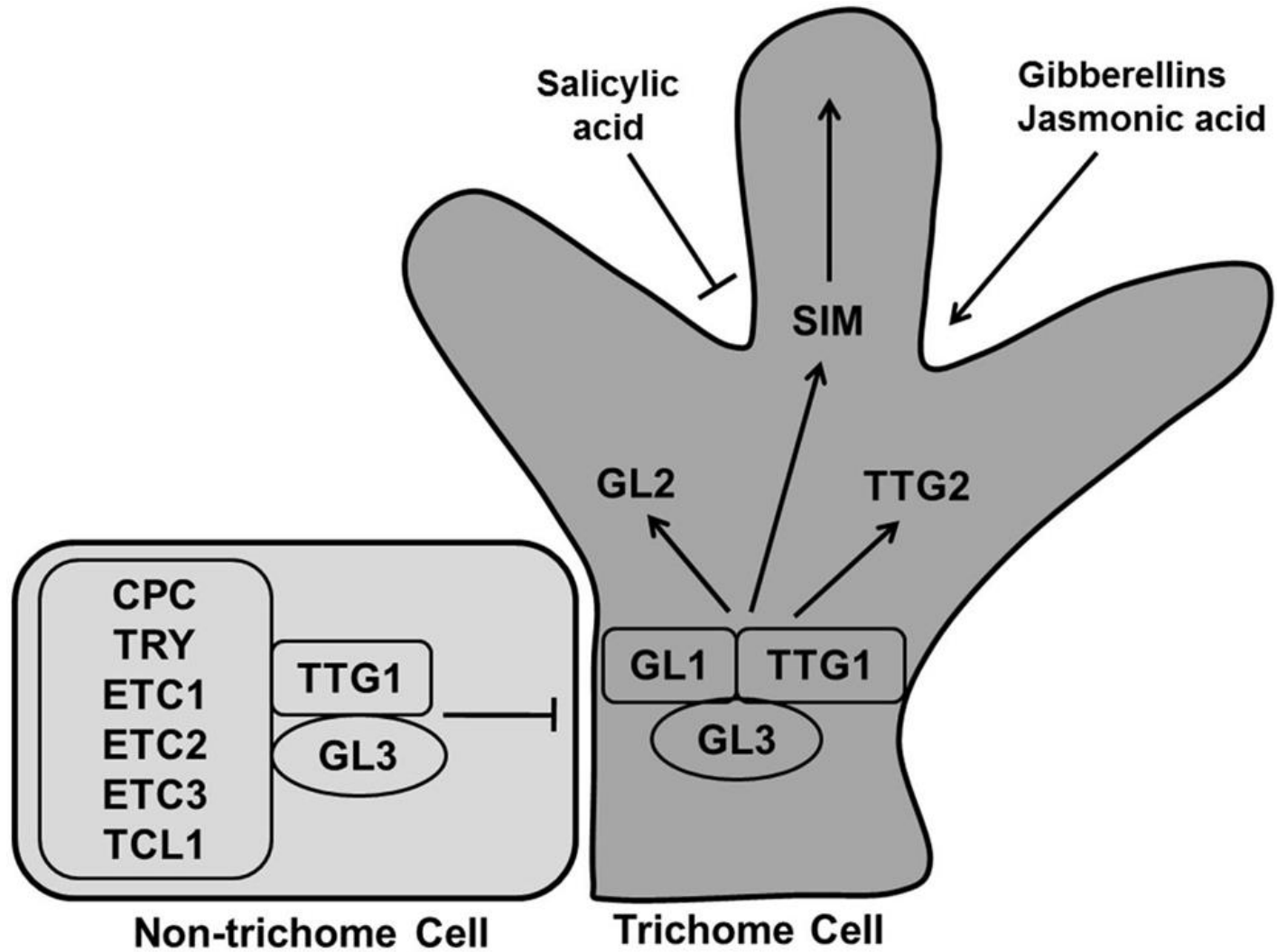


B. napus Trichome Genetics

Dos2 X NAM-0 F2 Doubled Haploid Phenotype



Trichome Regulation



B. villosa Interspecific Crosses

Hairy *B. rapa*
Hairy *B. napus*



B. villosa



- Crosses between *B. rapa* (AA) and *B. villosa* (CC) to re-synthesize *B. napus* AACCC
- Crosses between *B. napus* (AACCC) and *B. villosa* (CC) to generate a triploid bridge line

B. villosa Interspecific Crosses

Glabrous *B. oleracea* (CC)



B. villosa (CC)



- Move hairy trait into a CC genome background more amenable to crossing
- Develop RIL population to map genes responsible for hairiness

***B. villosa* Trichome Genetics**

***B. villosa* x *B. oleracea* RIL F2 Phenotypes**



***B. villosa* Trichome Genetics**

***B. villosa* x *B. oleraceae* RIL F2 Phenotypes**





Summary

- Hairs/trichomes deter flea beetles
- Contemporary canola has a defective *GL3* gene
 - Repair with *A. thaliana*, *B. rapa* and *B. villosa* genes
 - Growth penalty unless *GL3* expression is fine-tuned
- Work to map and move hairy trait from *B. napus* and *B. villosa* into a canola-quality background



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