



# Steps towards understanding variation for water and nitrogen uptake and use in *Brassica napus*

Hossein Zakeri<sup>1</sup>, Lanette Ehman<sup>2</sup>, Tina Wambach<sup>1</sup>, Isobel Parkin<sup>1</sup>, Steve Robinson<sup>1</sup>, Eric Johnson<sup>1</sup>, Yantai Gan<sup>1</sup>, Raju Datla<sup>2</sup>, Jeff Schoenau<sup>3</sup>, Rosalind Bueckert<sup>4</sup>, Sally Vail<sup>1</sup>

Agriculture and Agri-Food Canada, Saskatoon Research Centre (1), National Research Council, Saskatoon (2)

Departments of Soil Science, University of Saskatchewan (3), Department of Plant Sciences, University of Saskatchewan (4)

## Introduction

Water and nitrogen (N) are the biggest limiting factors in cropping systems around the world. Along with agronomic practices, breeding for limited N and water availability will increase profits of crop production. Existing variation for N and water uptake and utilization, and for drought tolerance/avoidance mechanisms among *Brassicas* (1, 2) could be useful in improving canola for adaptability and sustainability. Identifying the underlying traits to high N use efficiency (NUE) and water use efficiency (WUE) could help select for highly productive cultivars under low N application and variable water availability.

## Objectives

- Identify crop and plant physiological and biochemical traits that are closely related to NUE, WUE and to drought tolerance/avoidance mechanisms in *B. napus*
- Develop high-throughput screening methods and protocols for identified traits
- Assess variation within primary and secondary gene pools of *B. napus* for the selected traits
- Phenotypically characterize germplasm resources such as the spring *B. napus* Nested Association Mapping population currently under development

## Preliminary results

### 1- NUE experiments

Canola genotypes were grown under low and high N availability in greenhouse and field, and compared for N uptake, N utilization and yield.

#### Average biomass, plant N, and yield of canola genotypes grown under two rates of N in the field and greenhouse

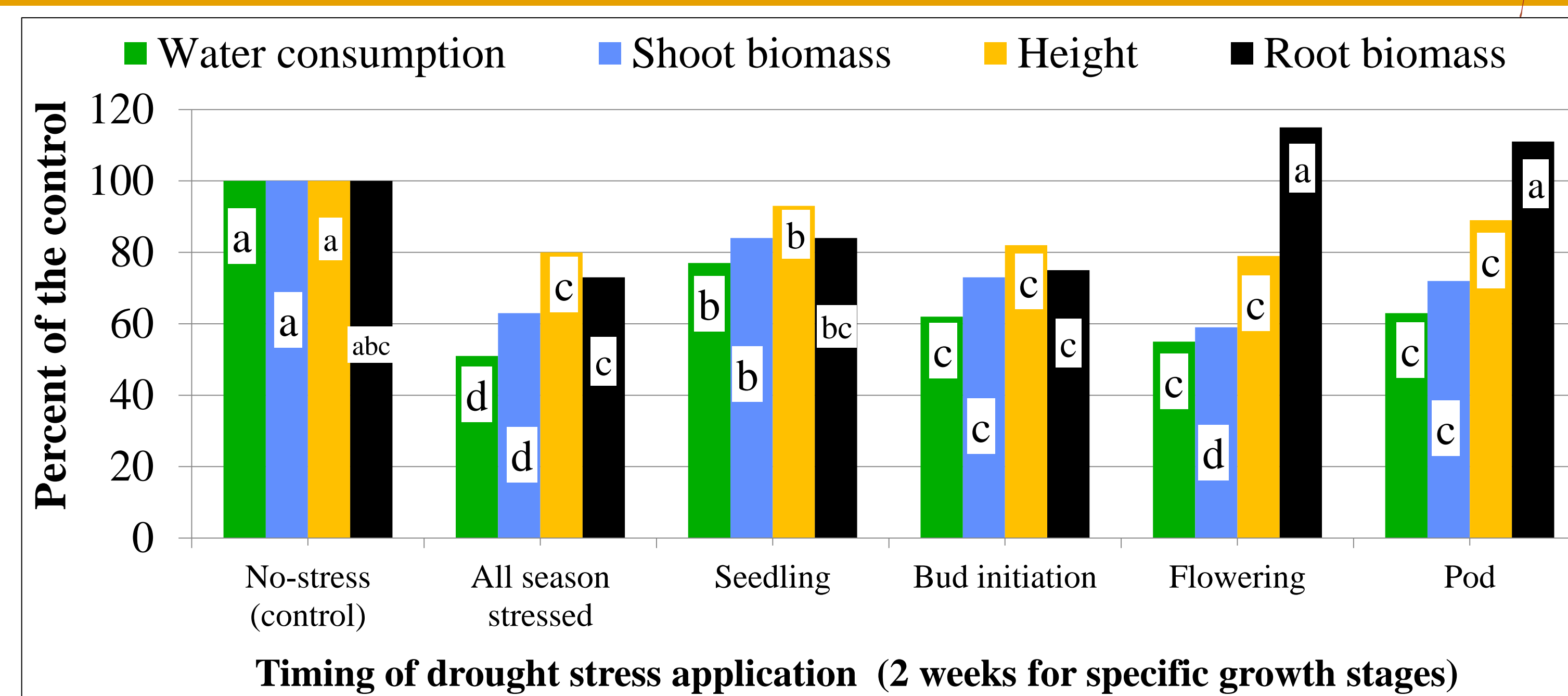
Soil N	Genotype	Greenhouse				Field location	
		Leaf N %	Stem N %	Plant N mg plant <sup>-1</sup>	Biomass <sup>†††</sup> g plant <sup>-1</sup>	Scott	Vanguard
Low N †	Hybrid	0.93 <sup>e</sup>	0.6 <sup>e</sup>	202 <sup>cd</sup>	15 <sup>ef</sup>	380 <sup>b</sup>	1551 <sup>a</sup>
	DH12075	0.9 <sup>e</sup>	0.9 <sup>cd</sup>	71 <sup>e</sup>	12 <sup>f</sup>	340 <sup>b</sup>	608 <sup>bc</sup>
	TIP2 (2)	1.0 <sup>de</sup>	0.7 <sup>de</sup>	187 <sup>cd</sup>	11 <sup>f</sup>	333 <sup>b</sup>	382 <sup>d</sup>
	N00-C171	1.0 <sup>de</sup>	0.8 <sup>de</sup>	140 <sup>de</sup>	12 <sup>f</sup>	--	--
	N99-508	1.2 <sup>cde</sup>	0.7 <sup>de</sup>	150 <sup>de</sup>	12 <sup>f</sup>	--	--
	YN01-429	0.85 <sup>e</sup>	0.5 <sup>e</sup>	141 <sup>de</sup>	13 <sup>f</sup>	--	--
High N ††	Hybrid <sup>a</sup>	1.7 <sup>bc</sup>	0.8 <sup>de</sup>	420 <sup>b</sup>	27 <sup>ab</sup>	513 <sup>a</sup>	1600 <sup>a</sup>
	DH12075	1.5 <sup>bcd</sup>	1.2 <sup>abc</sup>	241 <sup>c</sup>	29 <sup>a</sup>	603 <sup>a</sup>	838 <sup>b</sup>
	TIP2 (2)	1.9 <sup>ab</sup>	1.3 <sup>a</sup>	614 <sup>a</sup>	22 <sup>bcd</sup>	559 <sup>a</sup>	532 <sup>c</sup>
	N00-C171	2.4 <sup>a</sup>	1.3 <sup>ab</sup>	420 <sup>b</sup>	18 <sup>de</sup>	--	--
	N99-508	2.4 <sup>a</sup>	0.9 <sup>cd</sup>	395 <sup>b</sup>	21 <sup>cd</sup>	--	--
	YN01-429	1.9 <sup>ab</sup>	0.7 <sup>de</sup>	412 <sup>b</sup>	23 <sup>bc</sup>	--	--

Means within a column followed by the same letter are not significantly different ( $P < 0.05$ ). † Low N was 150 mg N pot<sup>-1</sup> in the greenhouse and 0 Kg N ha<sup>-1</sup> in the field - †† High N was 750 mg N pot<sup>-1</sup> in the greenhouse and 160 kg N ha<sup>-1</sup> in the field - ††† Above ground biomass

- ✓ Variation for total plant N content and N partitioning among leaf, stem and pod were identified among the genotypes under high N only
- ✓ Days to flowering and maturity affected plant N accumulation and partitioning
- ✓ Canopy closure, leaf area duration, leaf chlorophyll content, and N remobilization will be included in future studies

### 2- WUE experiment

Eleven canola genotypes were grown under six watering regimes: i) Fully watered as control (80% of the soil water holding capacity), ii) low water treatment (50% of soil water holding capacity during plant life cycle), two weeks water stress starting from iii) seedling, iv) bud initiation, v) flowering, vi) pod filling. Two weeks stressed plants received 30% of the soil water holding capacity. Plants were watered every other day.



Means of each trait (colour coded in legend) followed by the same letter are not significantly different ( $P < 0.05$ ).

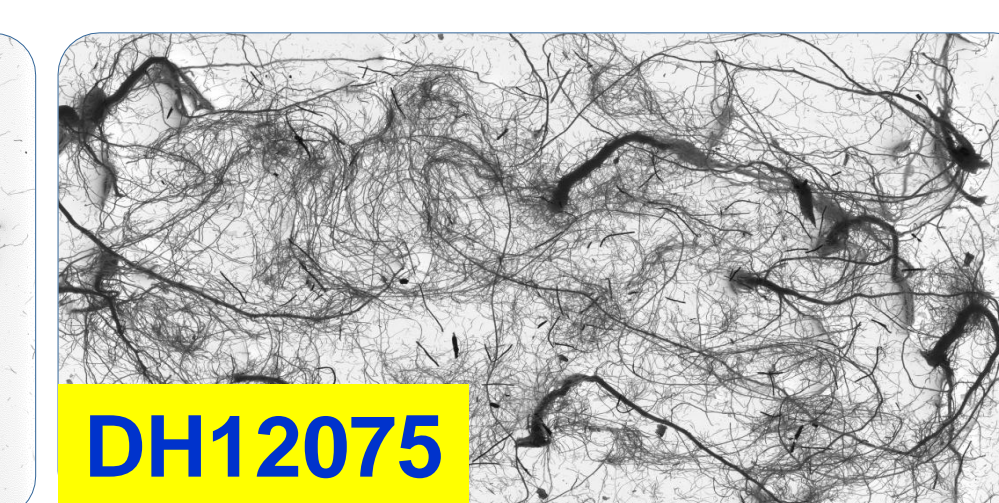
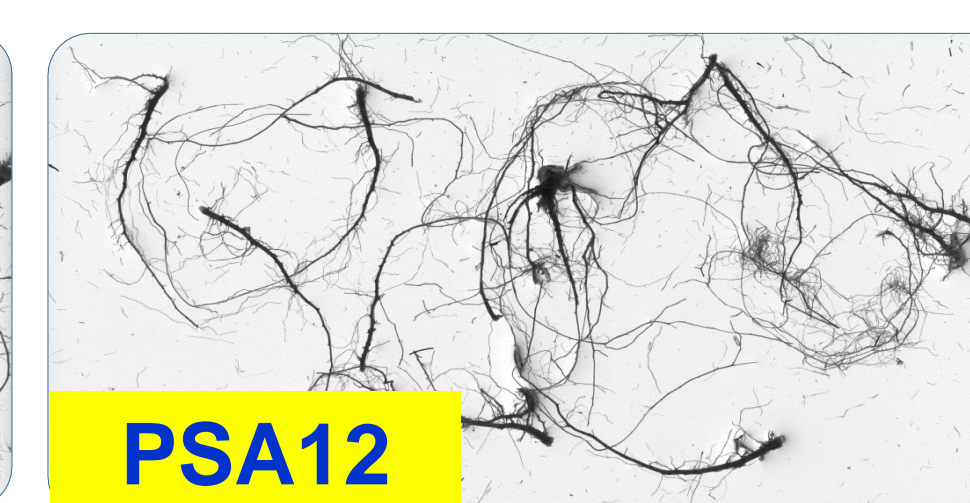
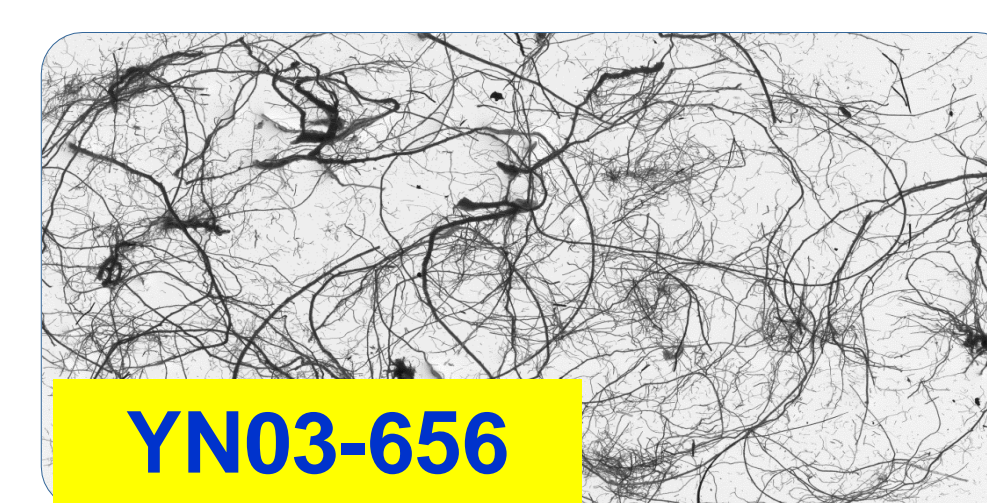
#### Above: Response of the genotypes to the watering treatments

#### Below: Variation of the genotypes, averaged over the water treatments

Genotype	Total water use	Plant height	Shoot biomass	Root mass	SPAD readings*
	(cm <sup>3</sup> plant <sup>-1</sup> )	cm	(g plant <sup>-1</sup> )	(g plant <sup>-1</sup> )	--
86004	11830 b	113 cd	26 ab	0.78 a	48 bcd
81N064 -5	7940 d	106 d	14 gh	0.23 gh	52 a
DH12075	14013 a	124 ab	28 a	0.61 b	48 bcd
DH38060	9609 c	106 d	18 ef	0.31 feg	50 bc
Karat	11788 b	110 d	22 cd	0.52 bc	45 de
N00 – C125	11414 b	120 bc	24 bc	0.41 cde	47 de
N99 – 508	10653 b	106 d	18 ef	0.34 def	49 bcd
PSA12	5720 e	109 d	12 h	0.14 h	55 a
Yickadee	10646 bc	108 d	20 de	0.45 cd	52 ab
YN03-656	13355 a	131 a	26 ab	0.39 def	48 bcd
YN04-C1213sp09	10194 c	126 ab	19 ef	0.29 efg	44 e

\* SPAD readings, recorded prior to flowering, reflects leaf chlorophyll content  
Means within a column followed by the same letter are not significantly different ( $P < 0.05$ ).

- ✓ Two weeks of water stress substantially reduced plant water consumption; whereas biomass reduction due to the stress varied by the treatment
- ✓ Genotypes substantially varied for total water consumption, for water consumption after two weeks stress, and for rooting systems
- ✓ Stress at seedling had the least effects on plant biomass and stress at flowering had the greatest negative effect among the stressed treatments
- ✓ Mechanisms of drought tolerance depended on the time of water shortage. Plants recovered shoot-biomass growth after the seedling stress, but had reduced height and extended roots due to the stress after flowering
- ✓ DH12075 consumed the most water and produced one of the largest biomass, whereas PSA12 had the lowest water usage and root mass



#### Root images of three genotypes of fully watered plants in the greenhouse

## References

- 1- Gan et al. 2004. Can. J. Plant Sci. 84: 697–704.
- 2- Svečnjak and Rengel. 2004. Field Crop Res. J. 97: 221-226.

## Acknowledgements

Technical help from R. Vetter, B. Hope, D. Williams, C. Headley, K. Livingstone, T. Mayama  
Funding from AAFC, NRC, ADF, SCDC.