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# Resistant sources and resources for clubroot at AAFC, Saskatoon

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Canada 

# Resistance to clubroot in *Brassica* species

**Black mustard; rich**

Brassica nigra

n=8

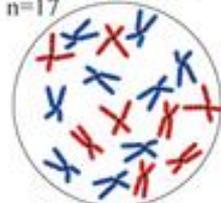


BB

Biofuel; no

Brassica carinata

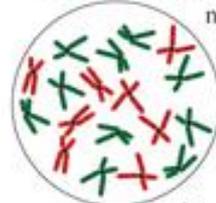
n=17



BBCC

Brassica juncea

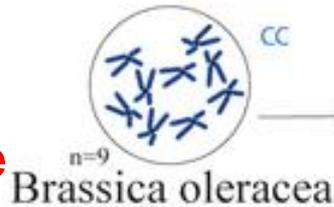
n=18



AABB

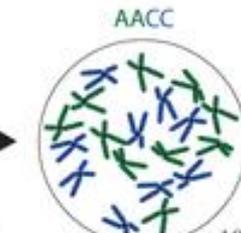
Mustard & canola;  
no

Vegetables;  
weak resistance



CC

Brassica oleracea



AACC

Brassica napus

Canola: no

Rutabaga: yes



AA

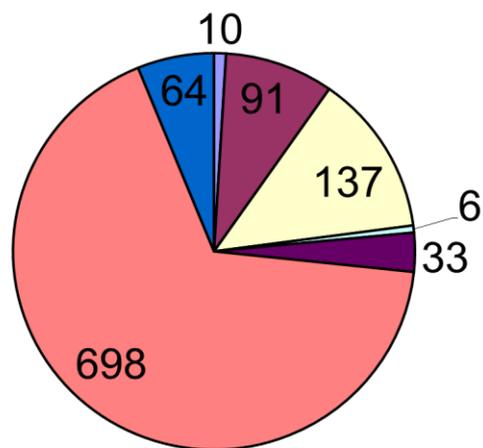
Brassica rapa

Vegetables;  
strong resistance  
originating from  
turnip

# Identifying sources of resistance at AAFC

• *P. brassicae* pathotype 3

■ *B. carinata* ■ *B. juncea* ■ *B. napus* ■ *B. nigra* ■ *B. oleracea* ■ *B. rapa* ■ Other



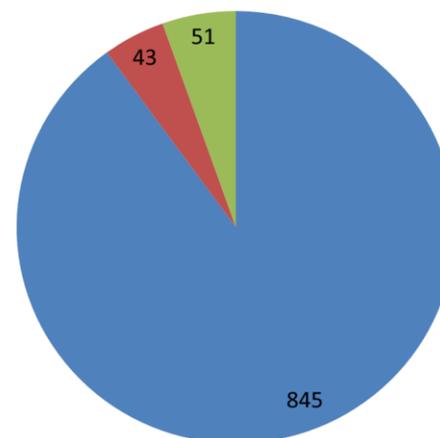
No. of R lines identified

<i>B. rapa</i>	24
<i>B. oleracea</i>	2
<i>B. nigra</i>	3
<i>B. napus</i>	0

Peng G, K.C. Falk et al 2014. Can J Plant Pathol, 36: 89-99.

• *P. brassicae* pathotype 5X

■ *B. napus* ■ *B. oleracea* ■ *B. nigra*



No. of R lines

<i>B. napus</i>	20
<i>B. oleracea</i>	2
<i>B. nigra</i>	7

Yu F et al, unpublished data

# Selected *B. napus* lines tested for pathotypes 3A, 2B and 3D

- 162 out of 845 *B. napus* lines were tested.
- Lines resistant to clubroot (0 DSI) were identified:
  - 3 resistant to all of the 4 pathotypes
  - 17 lines resistant to 1-3 pathotypes

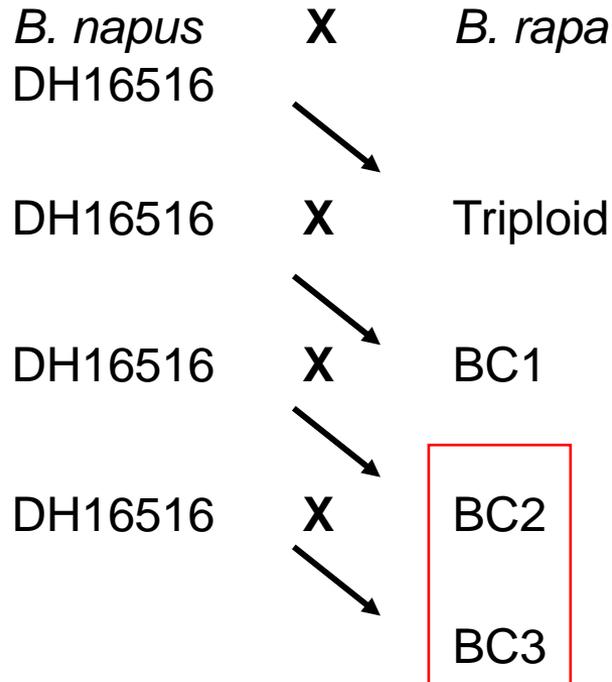
	2B	3D	5X-LG2
3A	0.77	0.50	0.31
2B		0.43	0.31
3D			0.29

# Nine *B. napus* lines were tested for resistance to more pathotypes

	2B	3A	3D	3H	3O	5C	5G	5K	5L	5X	8E	8J	8P
DH16516	S	S	S	S	S	S	S	S	S	S	S	S	S
Westar	S	S	S	S	S	S	S	S	S	S	S	S	S
AAFC682	S	S	S	S	S	R	R	R	S	R	S	R	R
AAFC693	R	S	S	S	S	R	R	R	S	R	S	R	S
AAFC694	S	S	S	S	S	R	R	R	S	R	S	R	S
AAFC787	S	S	S	S	S	S	R	R	S	R	S	R	S
AAFC692	S	R	R	R	S	R	R	R	R	R	R	R	R
AAFC405	S	R	R	R	S	R	S	S	R	R	R	R	R
AAFC660	R	R	R	R	R	R	R	R	R	R	R	R	R
AAF 665	R	R	R	R	R	R	R	R	R	R	R	R	R
AAFC695	R	R	R	R	R	R	R	R	R	R	R	R	R

➤ Three *B. napus* lines resistant to all of the pathotypes were identified.

# Introgression of clubroot resistance from turnip into canola



Microspore culture for developing doubled haploid (DH) *B. napus*



## Developing DH *B. napus* lines carrying CR genes from turnip

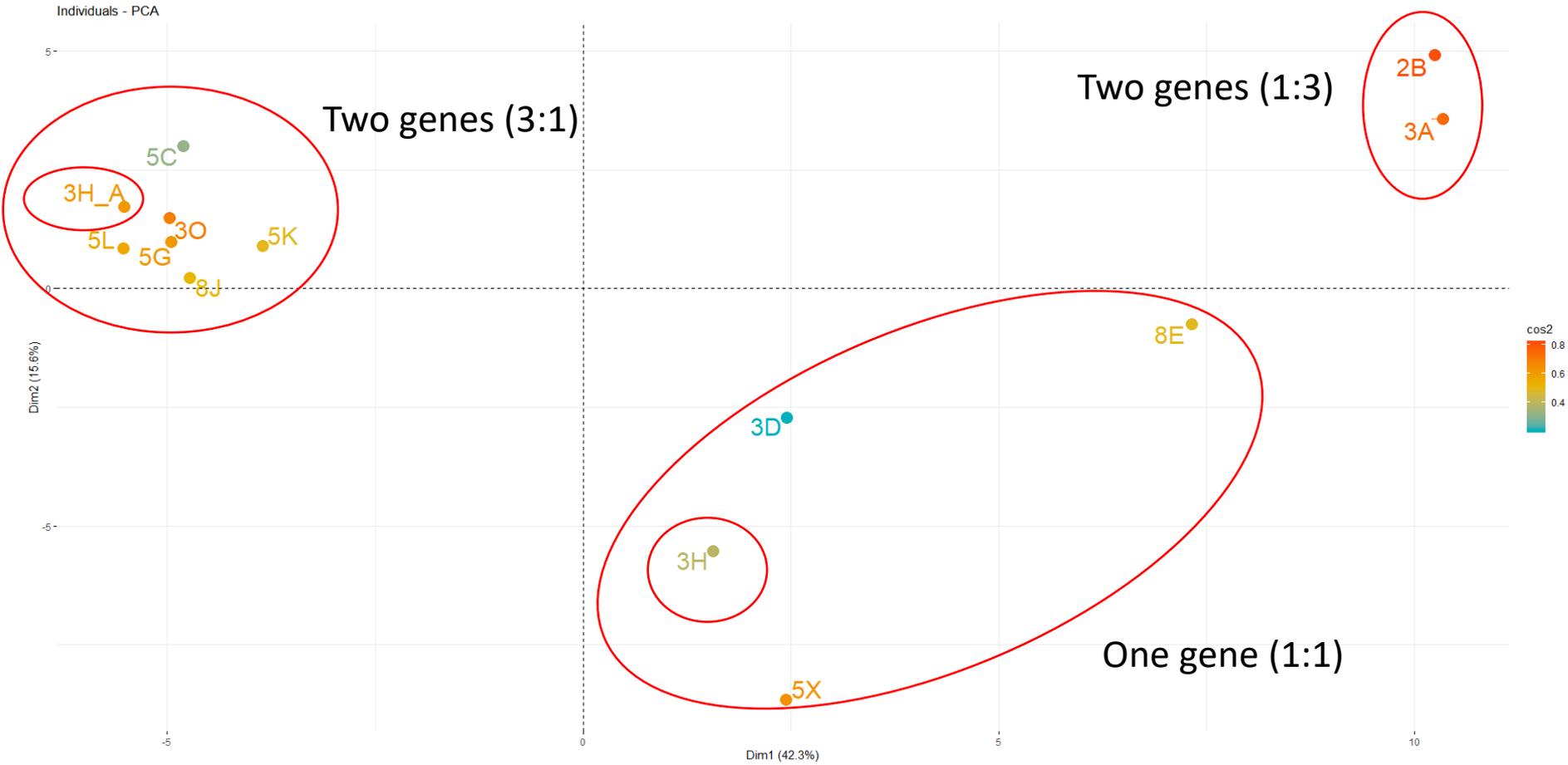
Donor Line	CR gene	Chr	Generation	No. of DH lines developed
Turnip Milan White	<i>Crr3</i>	A03	BC3	150
ECD01	<i>CRb</i>	A03	BC3	100
ECD02	<i>CRa</i>	A03	BC3	150
Debra	<i>CRc</i> <i>CRk</i>	A02 A03	BC2/BC3	190
Siloga	<i>Crr1</i> <i>Crr2</i> <i>Crr4</i>	A08 A01 A06	BC3	320

- Seed from > 800 DH lines was obtained.
- No vernalization is needed for all of the DH plants.
- Confirmation on the presence of each CR gene and identification of novel genes are in progress.

# Segregating for resistance and susceptibility in a DH pupation consisting of 84 DH lines from BC2 of DH16516 x Debra

Tested by	Pathoty pes	Average DSI	R	S	Expected ratio	<i>P</i>	No. of genes predicted
Stephen's group	3O	28.7	56	28	3:1	0.078	2
	5C	29.3	58	26	3:1	0.208	2
	5G	28.7	57	27	3:1	0.131	2
	5L	28.3	58	26	3:1	0.208	2
	5K	31.2	52	32	3:1	0.078	2
	8J	28.9	58	26	3:1	0.208	2
	3H_A	26.8	61	23	3:1	0.614	2
Fengqun's group	3H	34.2	51	33	1:1	0.050	1
	5X	38.7	49	35	1:1	0.127	1
	3D	34.5	48	36	1:1	0.190	1
	8E	47.4	37	47	1:1	0.275	1
	2B	54.3	25	59	1:3	0.313	2
	3A	55.6	27	57	1:3	0.131	2

# Relationship of resistance to different phenotypes among 84 DH lines



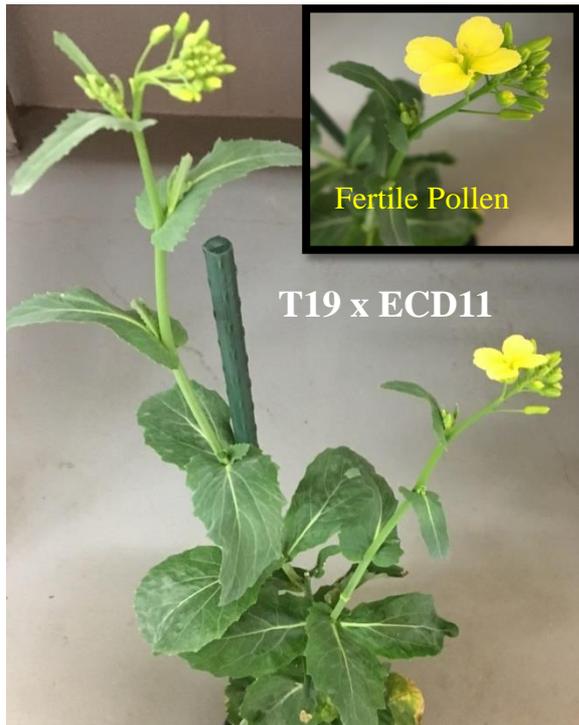


# Re-synthesizing *Brassica napus* with clubroot resistance from C-genome

**Yu F et al. 2017. Scientific Reports**

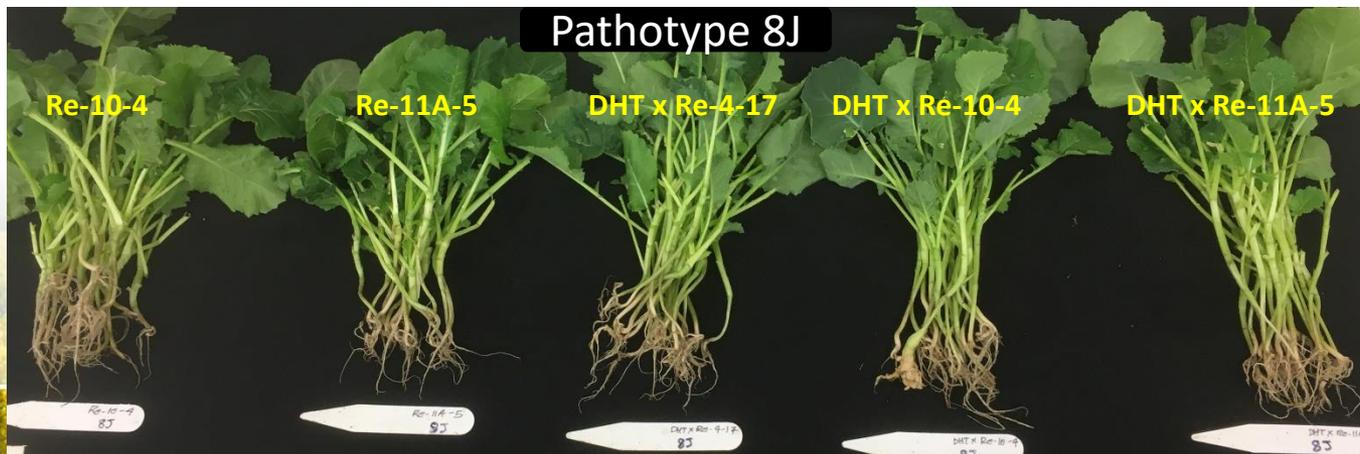


<i>B. rapa</i>	<i>B. oleracea</i>
T19 ( <i>Rcr4</i> , <i>Rcr8</i> and <i>Rcr9</i> )	ECD11
	JL04



# Evaluation of the re-synthesized *B. napus* lines from T19 x ECD11 for resistance to clubroot

Generation	Line name	3A_DSI(%)	8J_DSI(%)
F3	Re-10-4	41.0	0.0
F3	Re-11A-5	0.0	0.0
F1	DHT x Re-4-17	97.2	0.0
F1	DHT x Re-10-4	100.0	15.8
F1	DHT x Re-11A-5	100.0	0.0
S-control	DHT	100.0	100.0
R-control	45H29	100.0	88.9



# Summary on sources and resources for resistance Canadian pathotypes

- Sources of resistance: vegetables (*B. rapa*, *B. olerace* and *B. napus*) and mustard (*B. nigra*)
- More than 30 lines with good resistance were identified from the species at AAFC, Saskatoon.
- Resources were developed by introgression of resistance from some of the sources into canola/ *B. napus*.
- Genetic resources including SNP markers developed at AAFC have been / will be distributed to canola breeders for developing cultivars for resistance to clubroot.



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