Effects of Segment of 1RS of Rye on Root System of bread wheat Pavon Translocation 1RS.1BL

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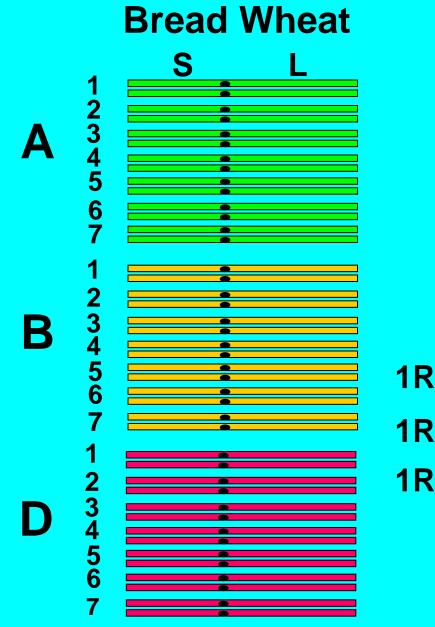
Ways Plants Survive Drought

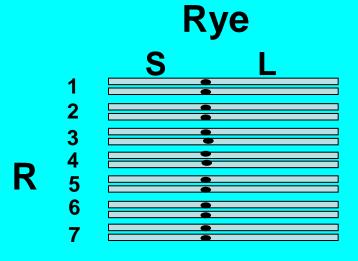
- Drought Escape

 crop earliness
- e
- Drought Resistance
 Dehydration Avoidance
 - maintenance of water uptake (Root System)
 - reduction of water loss
 - changes in tissue characteristics

Dehydration Tolerance

protoplasmic tolerance





Pavon Wheat











Why 1RS?

- Carried resistant genes to several pathogenes.
- On average, increased grain yield by 7% in certain cultivars.
- Increased adaptation and stability in stressful environments.
- No physiological basis for improved grain yield performance or better adaptability was given.

Pavon and its 1RS translocations







C ALE



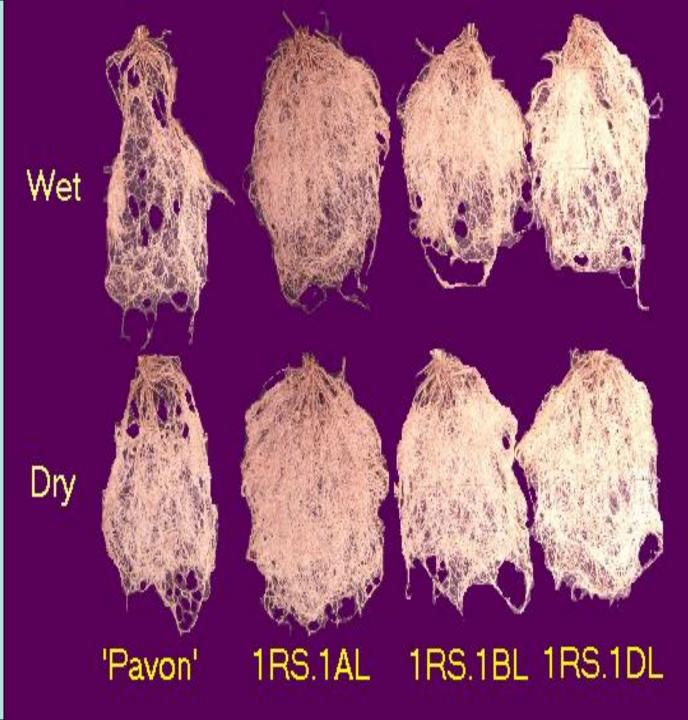


1RS : Short arm of chromosome 1 of Rye in Pavon wheat

Increase in Root Bio. 1RS.1AL 28% 1RS.1BL 11% 1RS.1DL 22%

Positive correlation between root biomass and grain yield in wet (0.66) and dry (0.89) pot conditions

Increase in GY in field 1RS.1AL 20% 1RS.1BL 11% 1RS.1DL 14%

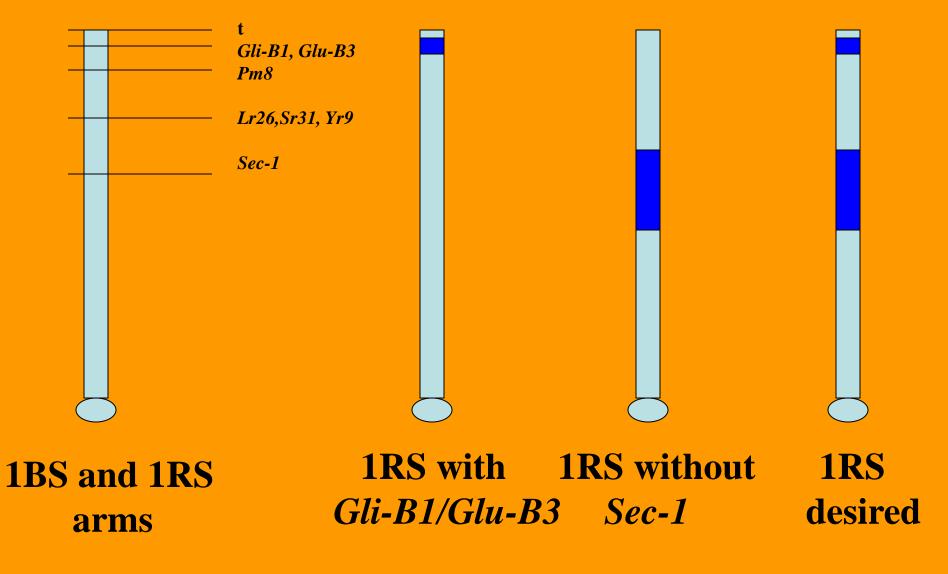


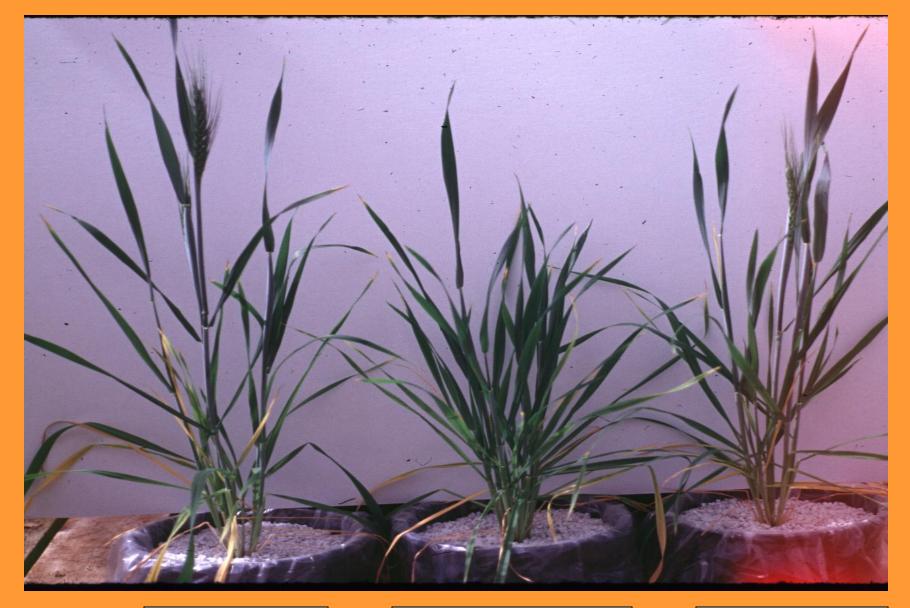
Disadvantage of 1RS

 Carries Sec-1 gene with negative effects on enduse quality.

- Lacks Gli-B1 and Glu-B3 genes with positive effects on end-use quality.
- 1RS.1BL lines are not used as commercial cultivars where bread quality is important.
- In China 50% and in Bulgaria 54% of the commercial bread wheat cultivars are of 1RS.1BL type.

Modified 1RS (segments of 1BS colored blue)







m1RS.1BL





Mean Plant Height (cm) Pavon 93 1RS.1BL 90 m1RS.1BL 78

Mean Days to Maturity

Pavon	121
1RS.1BL	126
m1RS.1BL	132

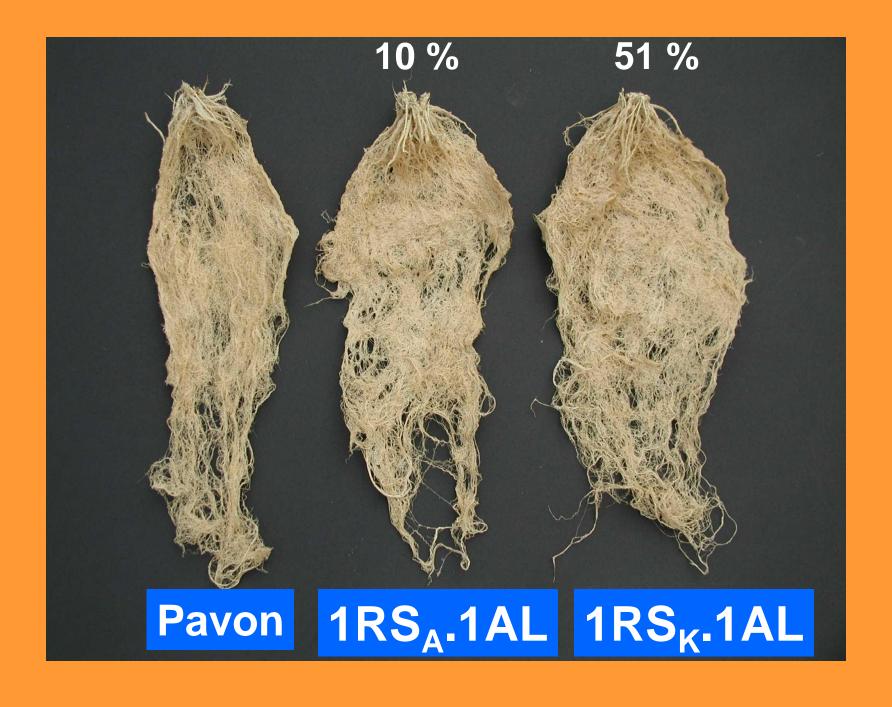


Root Biomass g/plant

Pavon 3.86 1RS.1BL 5.27 m1RS.1BL 5.26 Grain yield g/plant

Pavon 14.5 1RS.1BL 15.6 m1RS.1BL15.5





General Conclusions:

- Presence of 1RS in spring bread wheat increases root biomass in some 1RS.1BL.
- Location effect indicates interaction
 between wheat genetic background and 1RS
- Source of 1RS is important.
- Physiological basis of improved grain yield and adaptation may be due to larger root system in 1RS.1BL cultivars.

A larger root system would enhance water and nutrient uptake, which might contribute to increased shoot biomass and grain yield.

Wheat has two types of roots: Seminal roots, Nodal roots. The seminal roots are important for establishment of wheat seedling early in plant growth.

The nodal roots appear 4-5 weeks after germination and furnish most of the water and nutritional requirements of the plant under optimum conditions.

As water depleted from the tope layers of soil, seminal roots become more important to explore and furnish water from deeper soil layers. Deep seminal roots were associated with drought resistance through avoidance mechanism.

In CIMMYT wheats, 70% of total root length in found in the top 30 cm soil layer, where nutrient are concentrated in most agricultural soils.

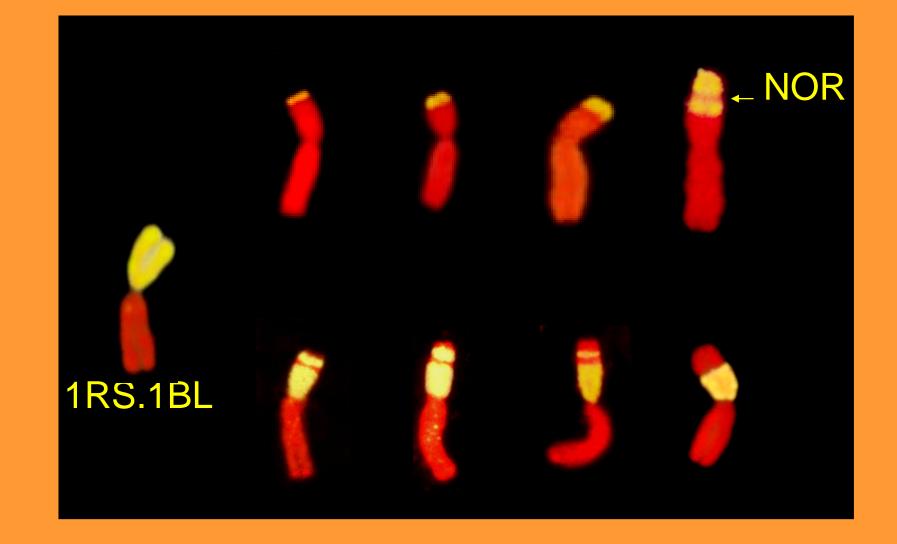
Pavon 1RS.1BL 1RS.1AL

14 Days After Sowing

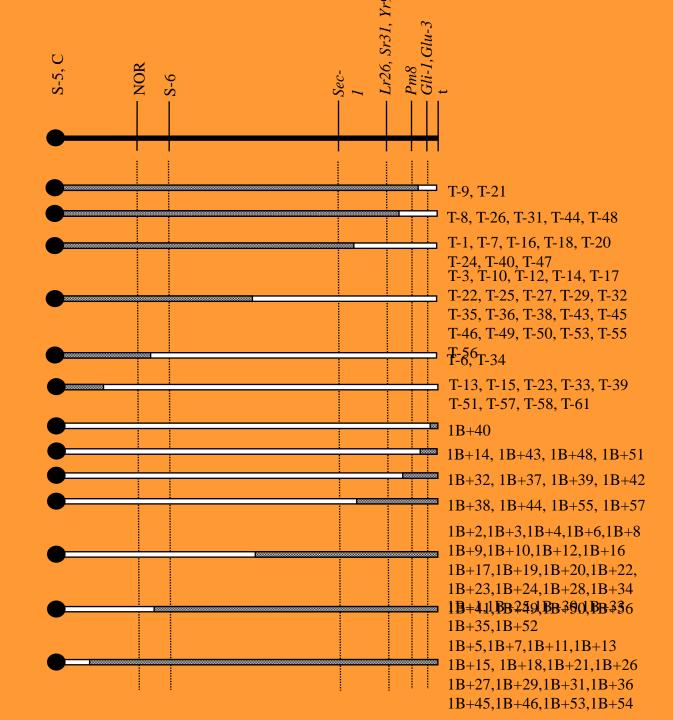
Pavon 1RS.1AL 1RS.1BL

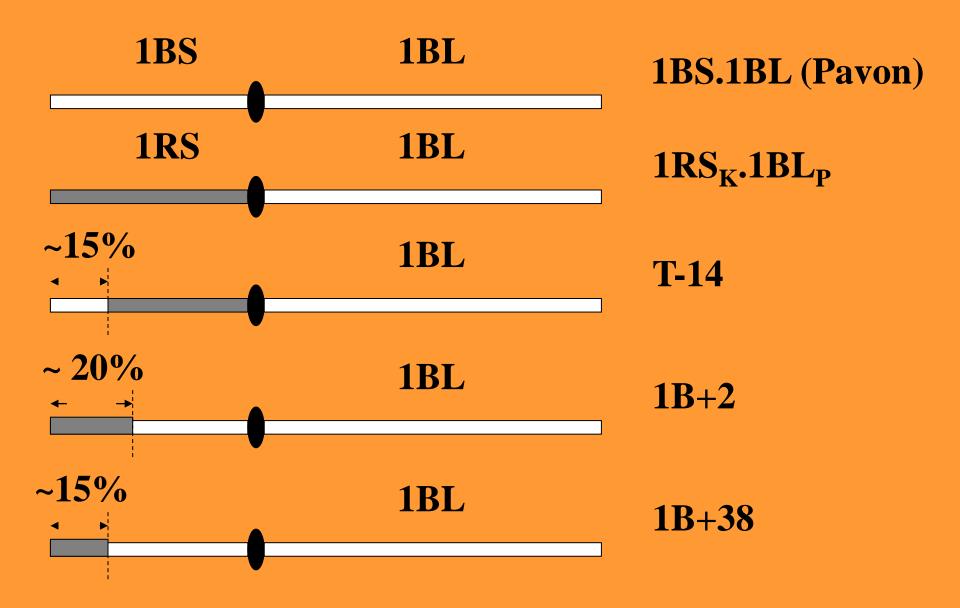
Questions:

- What segment (s) of 1RS carries gene (s) influencing root system?
- How 1RS affects root system?
- What is the genetic basis for 1RS variation?



The centric translocation 1RS.1BL and so recombinants of 1RS with wheat chromosome arm 1BS.





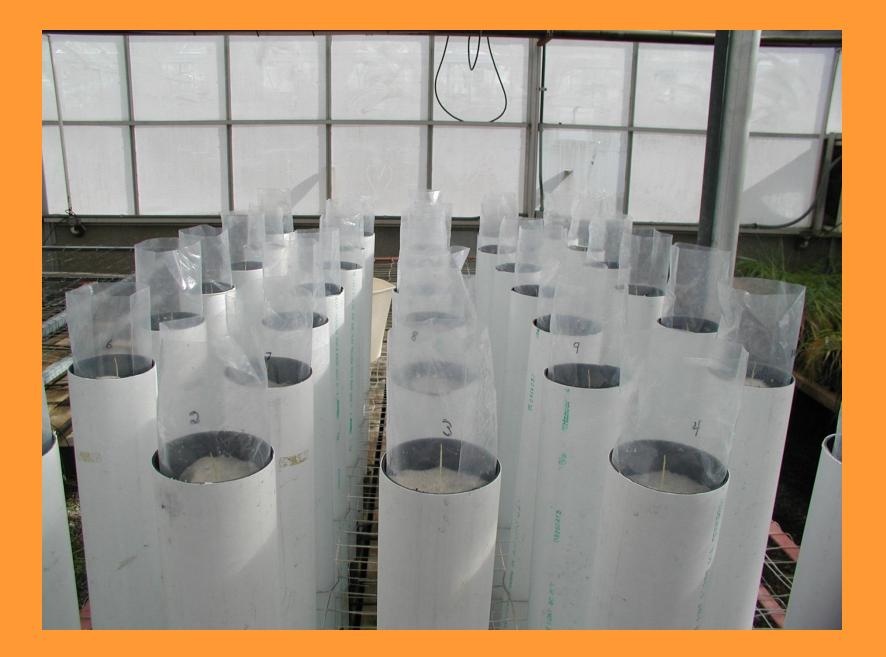
Tube experiment





















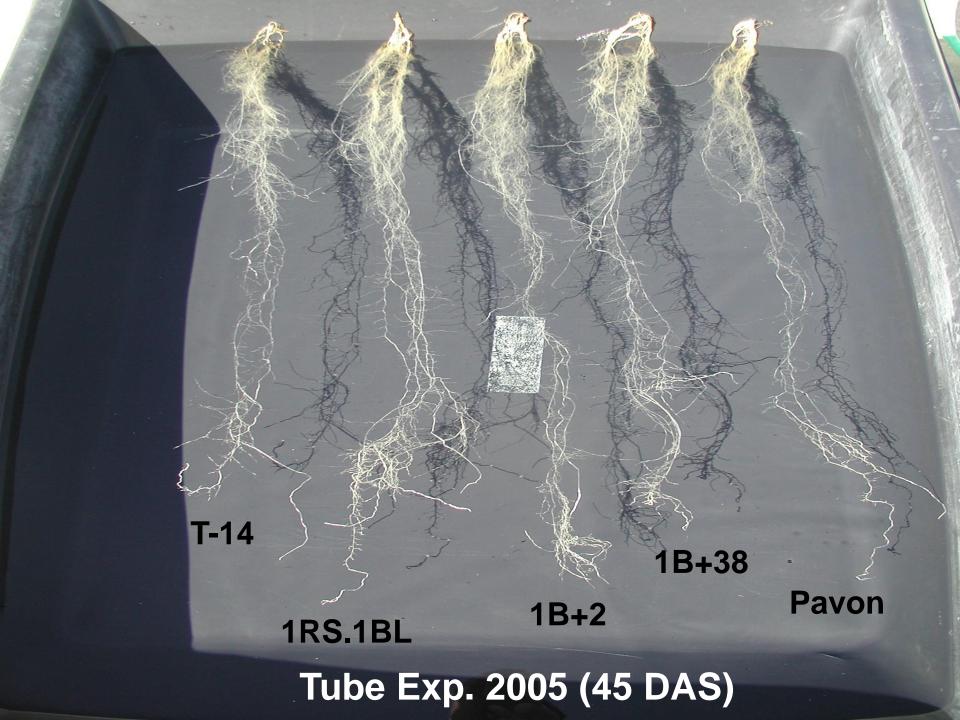


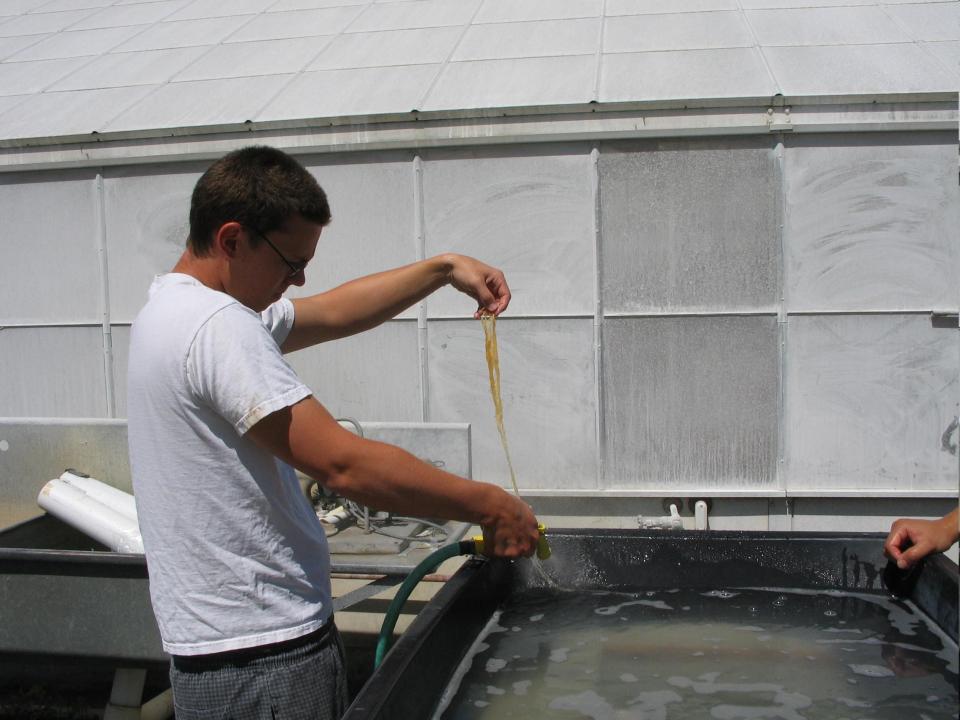






















	Plant height	No. of tillers	blade	Leaf blade length	Leaf blade area§		Shoot / root biomass
Genotype							
	cm	no.	mm	cm	cm ²	mg plant ⁻¹	mg/mg
Pavon	49.1 † a	3.1 ab	10.1 a	32.7 с	25.8 a	678.6 a	3.12 ab
1RS.1BL	51.0 a	3.3 ab	9.7 a	39.0 a	29.5 a	701.4 a	2.69 b
T-14	49.8 a	3.2 a	9.4 a	37.3 ab	27.4 a	557.1 a	3.16 a
1B+2	49.9 a	3.4 a	10.1 a	36.5 b	28.8 a	711.4 a	3.20 a
1B+38	49.6 a	3.0 b	9.9 a	37.1 b	28.7 a	681.4 a	3.14 a
CV‡(%)	5.7	10.0	9.4	6.1	14.2	17.5	12.99

§ Leaf blade area = leaf blade width \times leaf blade length \times 0.78.

Block 1

Genotype	NR>30	LRL	TRL	SRW	DRW	Rank
						sum
						(RAI)
G1	5 (<mark>2</mark>)	101 (1)	370 (1)	180 (<mark>2</mark>)	45 (<mark>1</mark>)	7
G2	6 <mark>(3</mark>)	103 (<mark>2</mark>)	470 (4)	195 (<mark>4</mark>)	70 (<mark>4</mark>)	17
G3	7 (4)	105 (<mark>3</mark>)	380 (<mark>2</mark>)	140 (1)	50 (<mark>2</mark>)	12
G4	4 (1)	110 (4)	420 (<u>3</u>)	185 (<mark>3</mark>)	55 (<mark>3</mark>)	14
Plack 2						

Block 2

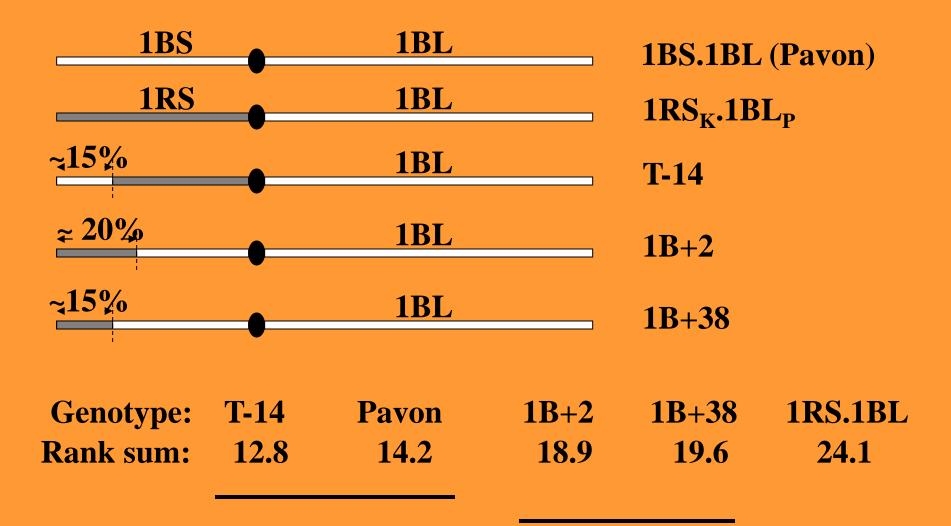
Block 3

ANOVA used to compare genotypes for each root traits.

To compare mean rank sums of the genotypes, rank sums in blocks were subjected to the statistical nonparametric Quade analysis developed for RCB designs. Quade test W is similar to F test in ANOVA.

Genotype	NR>30	LRL	TRL	SRW	DRW	RB	S _j §	RAI
	no.		-cm		mg			
Pavon	5.3† b	101 a	370 b	180 a	43 b	223 bc	-27.5	14.2 C
1RS.1BL	6.7 a	103 a	472 a	195 a	69 a	264 a	33.5	24.1 A
T-14	5.1 b	102 a	383 b	143 b	50 b	193 c	-26.5	12.8 C
1B+2	6.3 a	102 a	419 ab	185 a	54 ab	239 ab	1.5	18.9 B
1B+38	6.7 a	106 a	432 ab	168 ab	55 ab	223 bc	19.0	19.6 AB
CV‡ (%)	13.5	8	14	17	26	16		

§The critical value for a difference $|S_i - S_j|$ to be significant = 21.8



Conclusions:

- The distal segment of 1RS (~ 15% genetic map) carries major gene (s) enlarging root system in some 1RS.1BL bread wheat cultivars.
- There may be minor genes on other segments of 1RS affecting root system.
- Distal segments of chromosomes are subjected more to crossovers, therefore distal segments of different 1RSs have different genetic constitutions, and thus have different effects on root system and consequently on agronomic performance as reported in the literature.
- The distal segment of 1RS could be translocated to other wheat cultivars to manipulate their rooting ability, and thus improve their agronomic performance.
- Improved rooting ability should be coordinated with an appropriate preanthesis shoot system in order to accumulate maximum stem reserves as a consequence of greater water and nutrient uptake.

The End



Thank you for your listening.