



Breeding Strategies to Improve Nitrogen Use Efficiency in Oilseed Rape

J. Keurentjes, N. Erol-Oztolan, F. Breuer, A. Gertz, S. Miersch, H.C.Becker





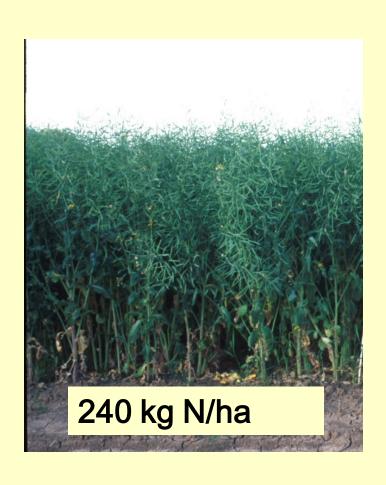
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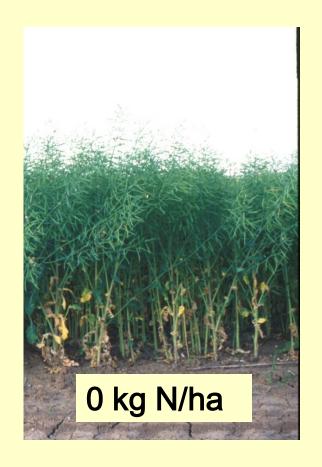
J. Keurentjes, N. Erol-Oztolan, F. Breuer, A. Gertz, S. Miersch, H.C.Becker

- Introduction
 - The problem
 - The NUE-CROPS research approach
- Arabidopsis
- Oilseed rape
- Conclusions and Outlook

The Problem

Oilseed rape reacts susceptible to reduced N supply



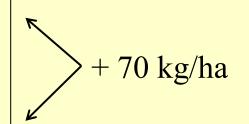


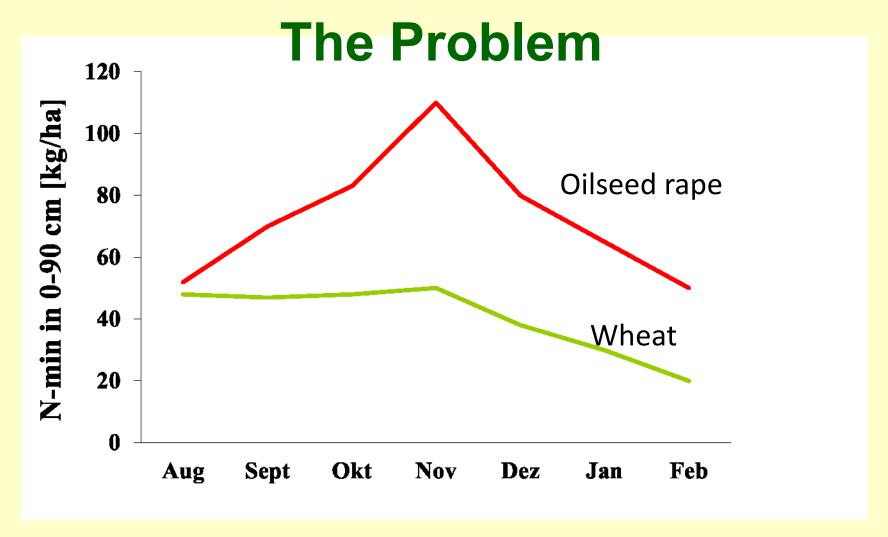
The Problem

- Oilseed rape is traditionally produced with high N-fertilizer inputs
- Oilseed rape is efficient in N uptake
- The N amount in harvested seeds is smaller than the N amount fertilized
- This may result in significant levels of nitrate leaching after harvest
- It is necessary to reduce the N surplus

Example	ļ
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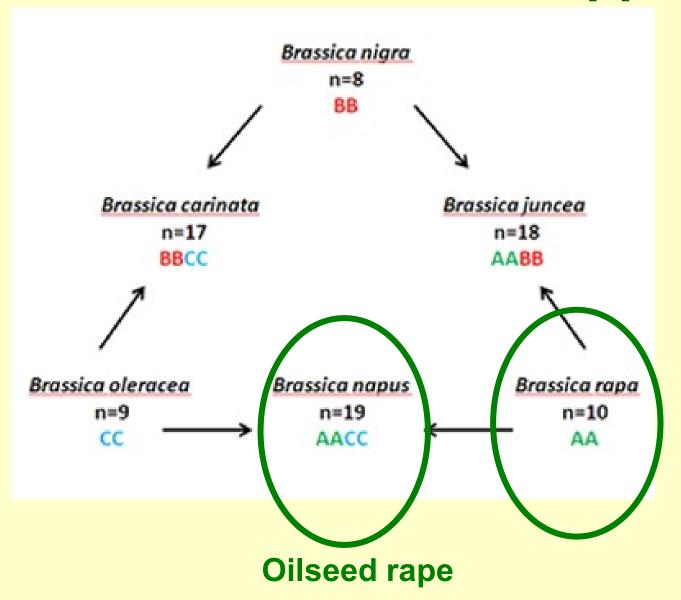
N Fertilization	180 kg/ha
N in total plant biomass	330 kg/ha
N in harvested seed	110 kg/ha

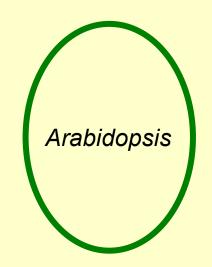




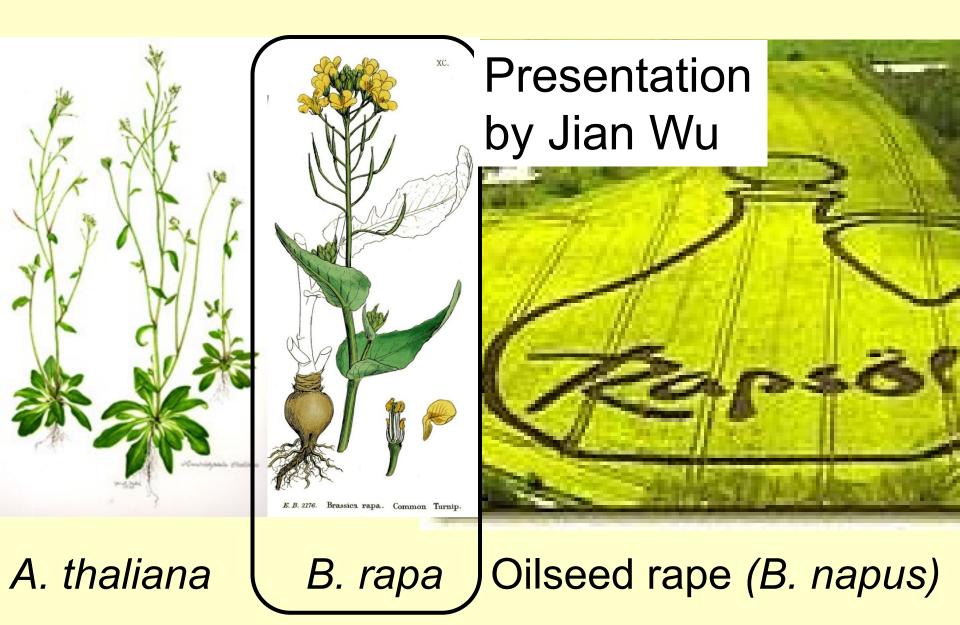
N-min after harvest of wheat and oilseed rape (2 years, Höckelheim, LK Hannover)

The NUE-CROPS approach





The NUE-CROPS approach



WP 3.1:

Genetic Analysis of Nitrogen Use Efficiency in *Arabidopsis thaliana*

Nihal Öztolan Erol, Joost Keurentjes





As a Model Organism *Arabidopsis* thaliana

- Dicot used as model organism in many studies
- 5 chromosomes, 120 mb genome and ~30,000 genes
- Brassicaceae family synteny
- Self fertilization
- World-wide natural variation



Materials and Methods

• Plant Material:

HapMap population (353 accessions)

• Growth Conditions:

- On rockwool bloks, covered to prevent algea
- 1 mM N (Deficiency) vs 5 mM N (Control)

Measurements After Harvesting:

- Shoot Fresh Weight (g), Shoot Dry Weight (g) (SW)
- Shoot N concentration (using DUMAS method)
- Nit_UI (SW(SW/N content))
- Nit_UE (SW/N content)

Association Mapping:

- Full set of densely SNP markers: 250,000 SNPs
- Based on Linkage Disequilibrium
- Therefore, detection of significant SNPs within 20kb will give us a QTL region which might be associated with NUE.

N deficiency

Control

a.



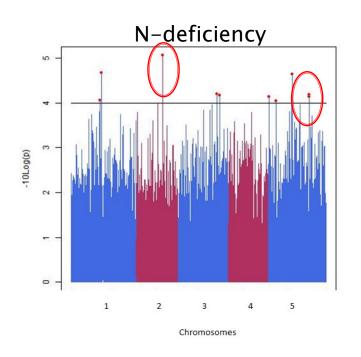


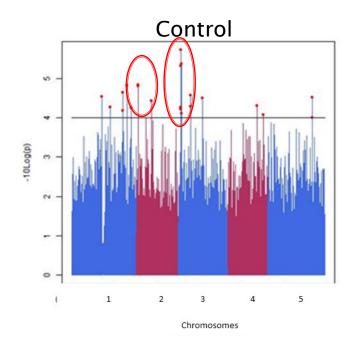
C.



Association Mapping

GWA Mapping Nitrogen Use Efficiency

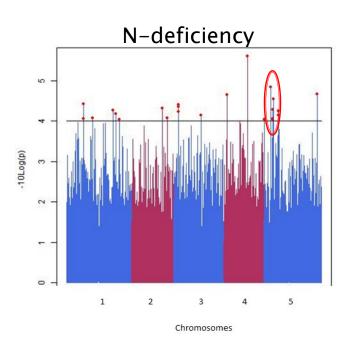




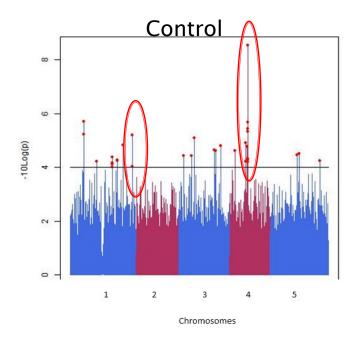
AT2G28800 Albino 3 (Alb3) AT5G47120 Bax inhibitor 1 (BI-1) AT5G47110 chlorophyll a/b binding protein gene (LIL3:2)

AT2G02590 Unknown AT3G05090 Lateral Root Stimulator 1 (LRS1)

GWA Mapping Dry Weight



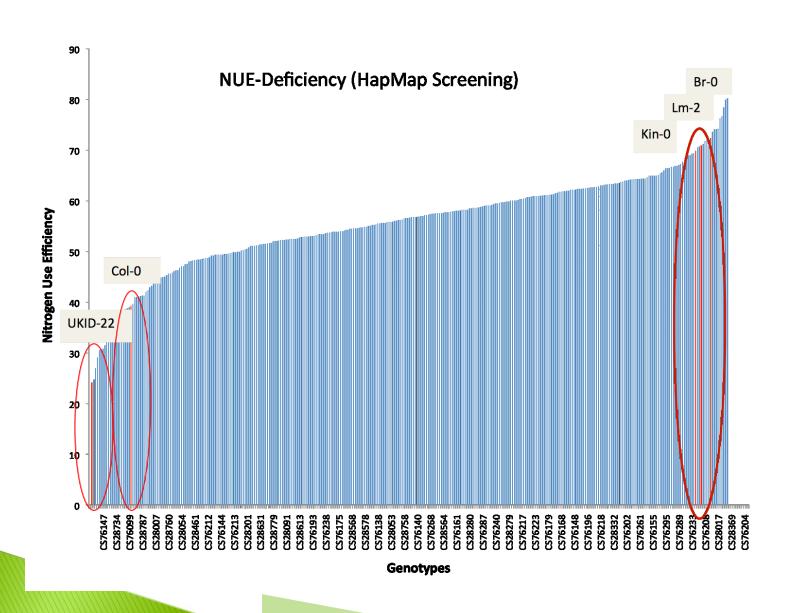
AT5G13550 Sulfate transporter (SULTR4;1)



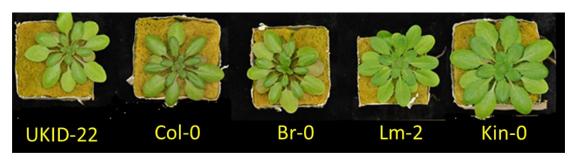
AT1G75960 AMP-dependent synthetase and ligase family protein AT4G14400 Accelerated cell death 6 (ACD6)

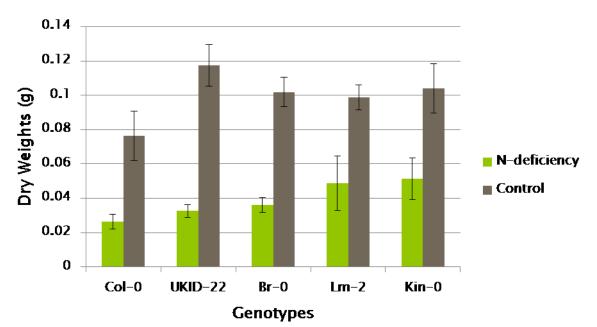
Growing Extreme Genotypes and Mutants

Ranking accessions









Summary *Arabidopsis*

Br-0 and Col-0 extreme genotypes

AT2G28800 Albino 3 (Alb3)

AT4G27010 Embryo defective 2788

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AT5G47120 Bax inhibitor 1 (BI-1)
AT5G47110 chlorophyll a/b binding protein gene (LIL3:2)
AT2G02590 Unknown
AT3G05090 Lateral Root Stimulator 1 (LRS1)
AT1G22170 Phosphoglycerate mutase family protein
AT2G14835 Ring/U box superfamily protein
AT5G13380 Auxin responsive GH3 family protein
AT4G14400 Accelerated cell death 6 (ACD6)
AT5G13550 Sulfate transporter (SULTR4;1)
AT1G75960 AMP-dependent synthetase and ligase family protein
AT1G52070 Mannose-binding lectine super family protein
AT3G46540 ENTH/VHS family protein
AT4G26990 Unknown
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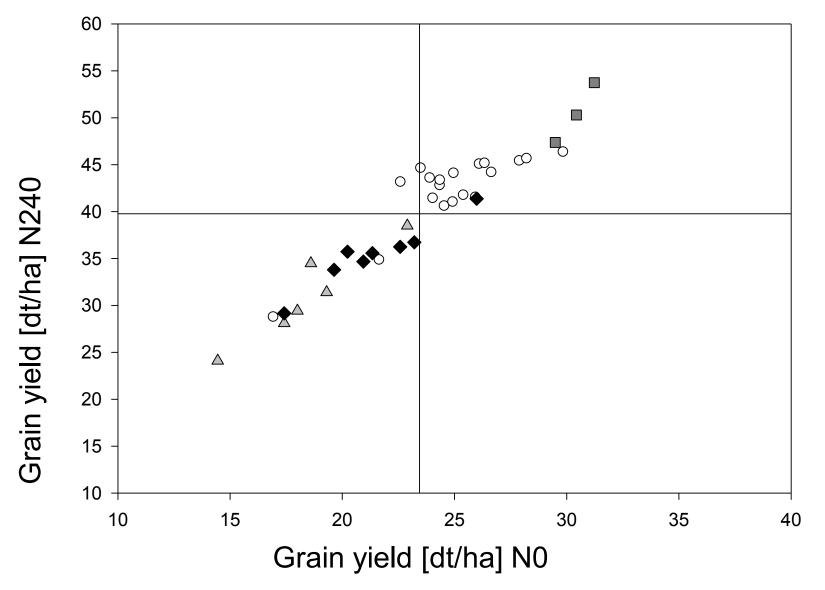






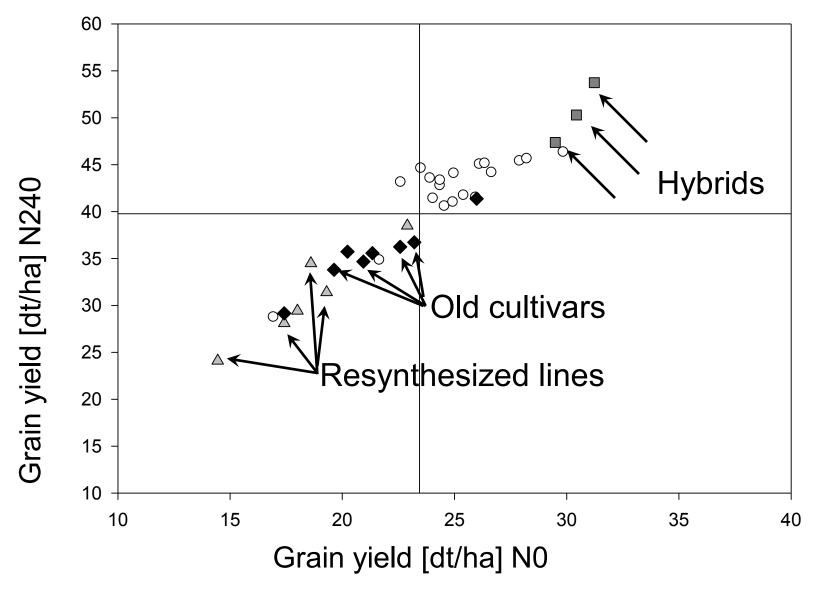
WP3.3 Oilseed rape

Selvastian Miersch, Frank Breuer, Andreas Gertz, Heiko Becker



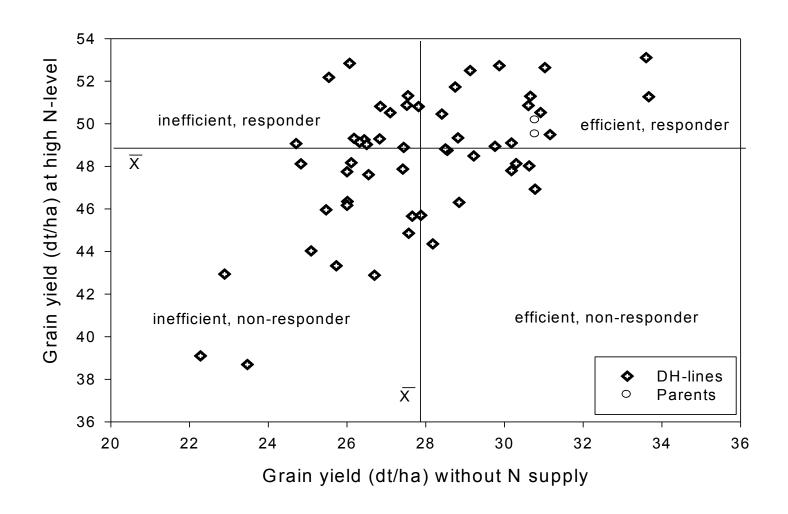
36 genotypes (resynthesized lines, old and new cultivars), 7 environments

Kessel et al. 2012



36 genotypes (resynthesized lines, old and new cultivars), 7 environments

Kessel et al. 2012



54 DH lines (Apex x Mohican), 7 environments

NYIKAKO et al. 2003

Variance components for grain yield of oilseed rape grown at two N levels

Source of variation	Broad material	DH Apex x Mohican	
	(n=36)	(n=56)	
Genotype	28.44	5.02	
Genotype x location	6.82	2.61	
Genotyp x N level	3.85	2.01	

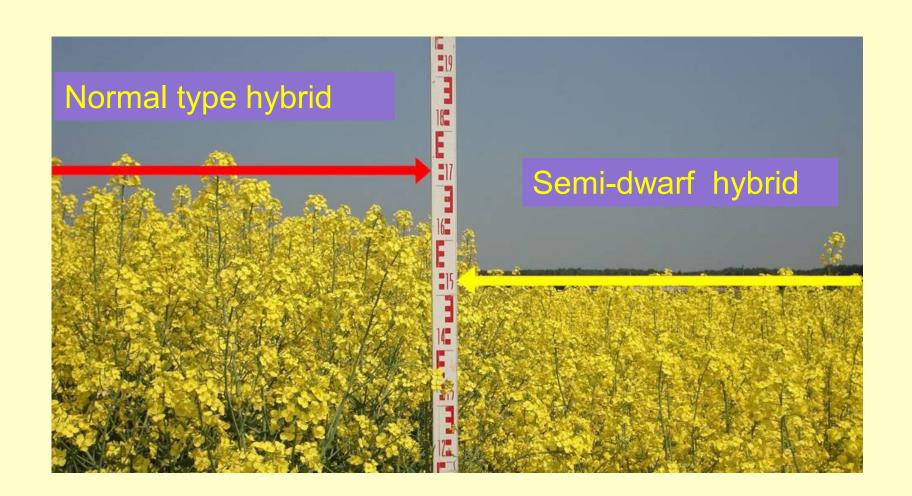
- → In unselected material there is a considerable variation in N efficiency
- → Large potential to select for N efficient cultivars

Are Semi-dwarf Hybrids a way to increase N efficiency?



Dwarf (ms mother)

Semi-dwarf Hybrid Normal type (pollinator)



Comparison of Semi-dwarf and normal hybrids

- > A dwarf line was crossed with a normal line
- F1 derived DH-lines (242 DH-lines) segregate into dwarf and normal types
- all DH Lines were crossed to a normal male sterile tester to produce normal and semi-dwarf testcross hybrids
- development of plant material by KWS Saat AG





Field trials 2011/12 and 2012/13

- 108 testcross hybrids (54 semi dwarf and 54 normal types)
 were sown in replicated field trials in Einbeck and Göttingen
- > Selection of genotypes based on differences in N uptake from 2010/11
- cultivated under low nitrogen supply (no fertilization) and optimal nitrogen supply (160 kg N/ha)
- two harvest dates:

end of flowering

maturity – 2011/12 hail in Einbeck – no grain harvest only straw







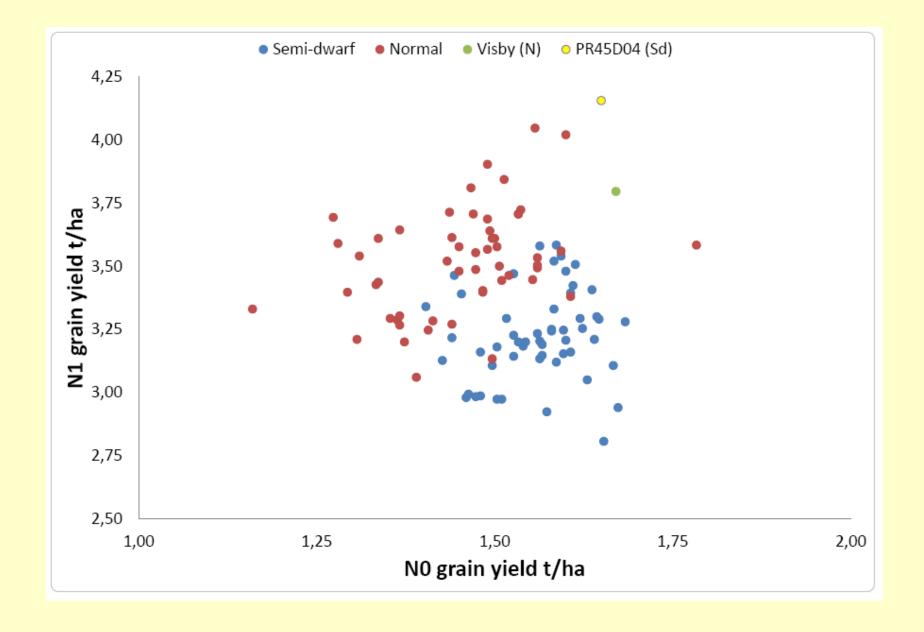
Linkage map

- A total of 881 (432 SNP, 109 SSR and 340 AFLP) polymorphic marker
- The final linkage map includes 438 markers (390 SNP, 49 SSR) distributed on 22 linkage groups (LG)
- 157 marker loci co-segregate
- 21 LGs could be assigned to the 19 chromosomes of the *Brassica napus* genome
- total map length of 1301 cM (Kosambi)

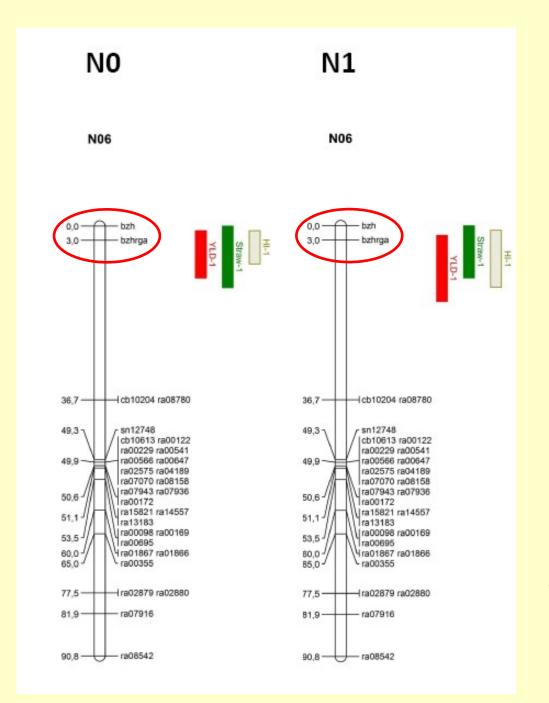
Results

Mean values of 54 semi-dwarf and 54 normal-type testcrosses (mean of 3 environments)

		Grain t/ha DM		Straw t/ha DM		н	
NO	Semi-dwarf	1.56	**	2.87	**	0.35	**
NO	Normal	1.45		3.48		0.29	
N1	Semi-dwarf	3.22	**	3.94	**	0.45	
IAT	Normal	3.52		4.36		0.45	



Yield of 54 semi-dwarf and 54 normal-type testcrosses (3 environments)



Major QTL for grain yield, straw yield and HI in the region of the dwarf gene

Conclusion and Outlook

- Nit-UE is a complex trait with considerable genetic variation
- Plant architecture (semi-dwarf) is of interest
- Role of root system is still largly unexplored
- Association mapping and QTL mapping are useful tools
- Combined analysis of all data from this sub-project is a challenge for the next few months

Root system size measured by electrical capacitance

(Chloupek 1977)



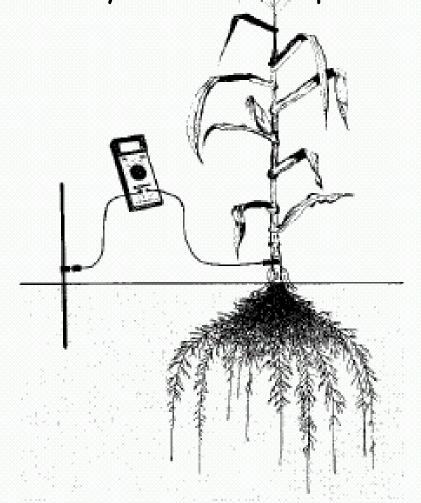


Photo: Julia RUDLOFF

Fig. 1. Estimation of root system size with a BK Precision 810A capacitance meter. The negative electrode is attached to the maize stem 6 cm above ground level and the positive electrode is attached to a copper ground rod.

Van Beem et al.

1998

