

The ABSTRESS project - Work Package 1: Define and establish the experimental conditions for investigating plant response to drought and Fusarium stress

Christophe Salon

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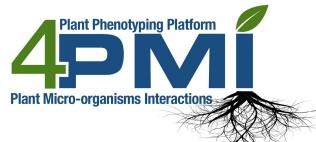






Work Package #1 : Define and establish une experimental conditions for investigating plant response to drought and Fusarium stress INRA, CSIC, FERA, ARTERA

Christophe Salon, Carmen Bianco, Adrian Charlton, Roberto Defez, Mike Dickinson, Rebecca Iglesias, Christian Jeudy, Tracy Lawson, Jack Mathews, Ulrike Mathesius, Phil Mullineaux, Marion Prudent, Nicolas Raspail, Diego Rubiales.









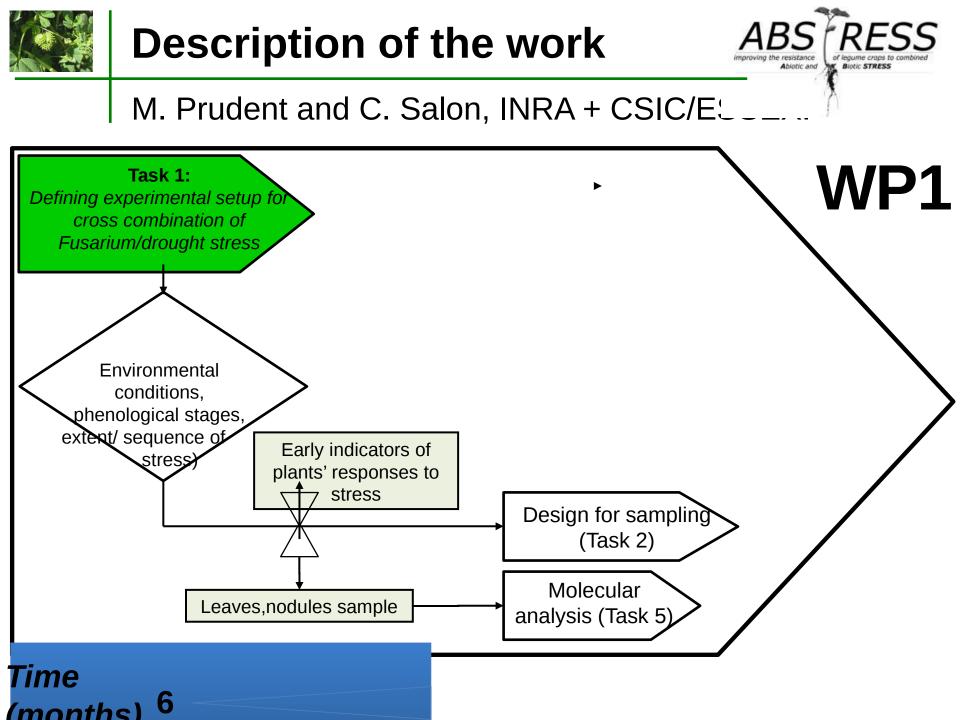
- **Protocols** for investigating resistance mechanisms to drought and Fusarium,
- Produce material for the consortium,
- **Characterize** plant/pathogen interactions under drought stress and compare with optimum environmental conditions in a factorial design,
- **Give guidance** to identified plant phenotypic traits for selecting genotypes having enhanced resistance to drought and Fusarium.

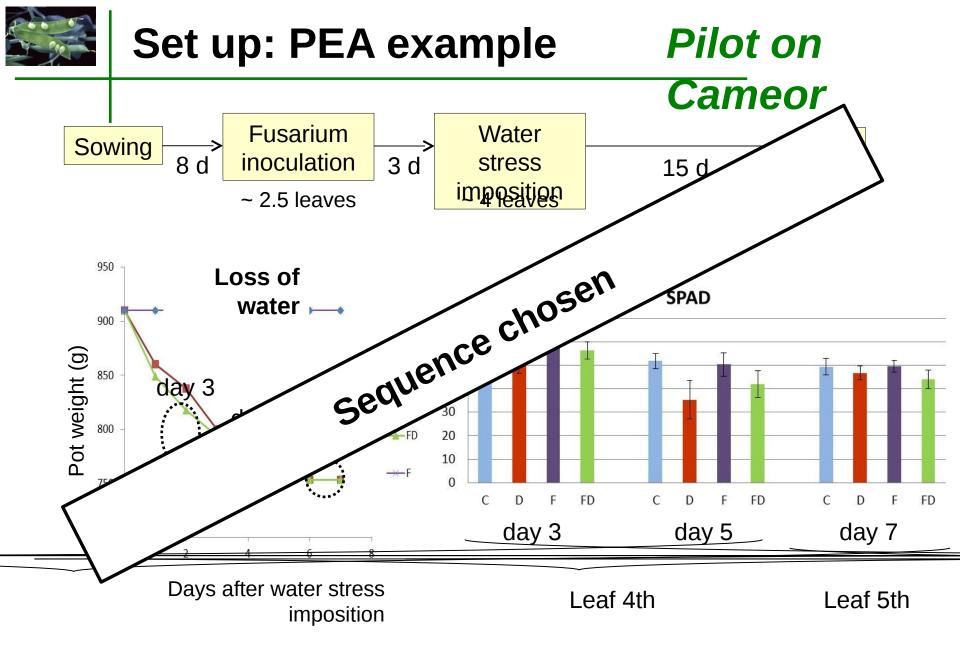


Iasks



Task No.	Activities	Inst. Responsible for Task	Person(s) Responsible for Task	
T1.1	Defining experimental setup for cross combination of <i>Fusarium</i> /drought stress	INRA/CSIC		





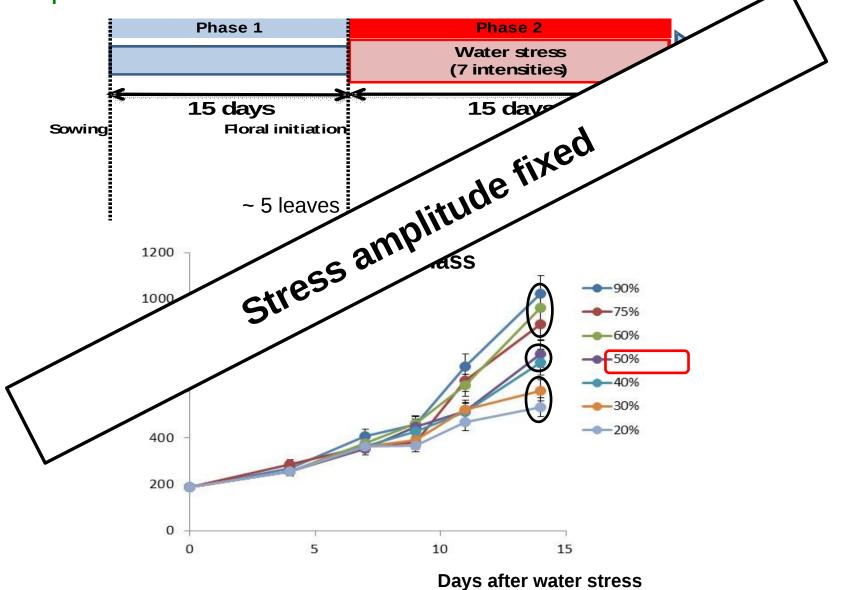
^{+ 3} volumes of Fusarium inoculation tested: 10, 20, 30, 50 mL



Set up: PEA example

Comparison of different water stress intertensity (12):

Stress

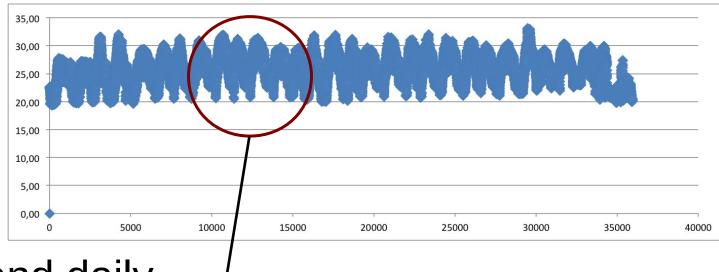


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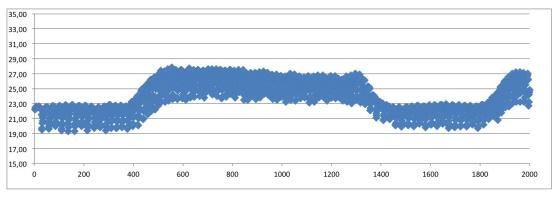
Set up: greenhouse conditions





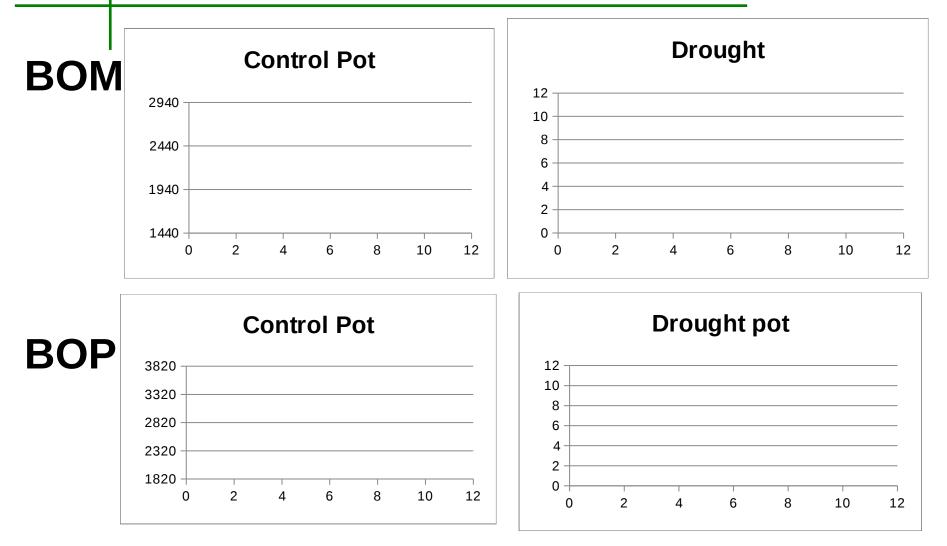
...and daily







Plant watering verified



Automatic

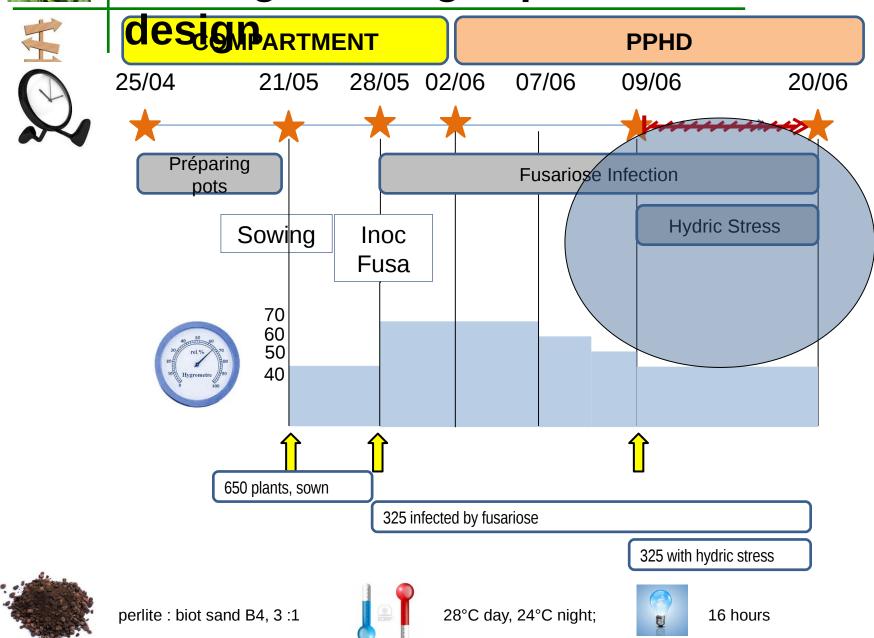


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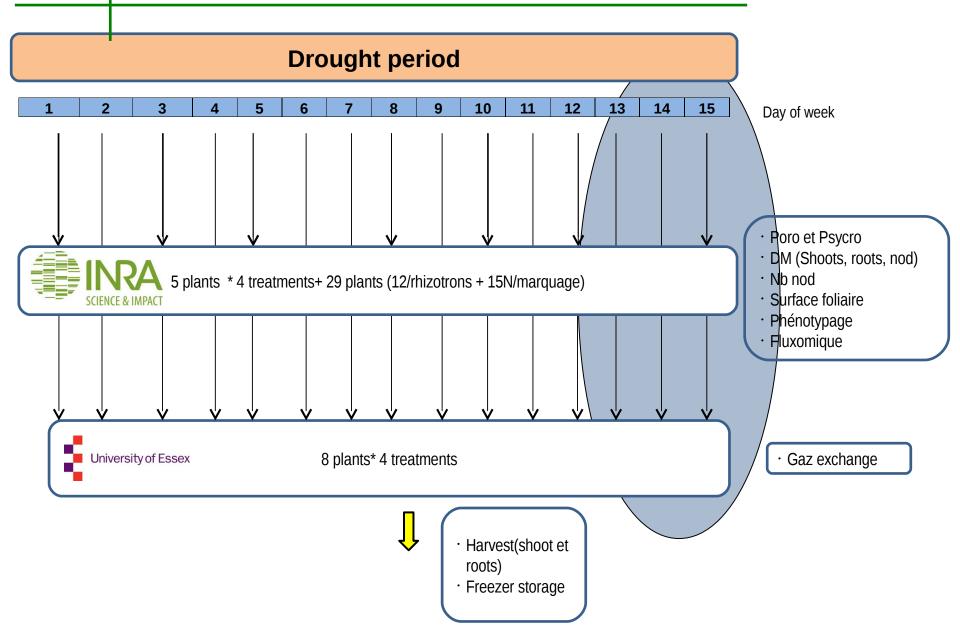


- Transferring protocols devised for the infection of plants with *Fusarium* in highly controlled environments: challenging in the local conditions in Dijon.
- Drought conditions, with fluctuations in the internal environment of the platform: challenging conditions in which to undertake controlled experiments.
- This added complexity for example the interpretation of physiological data
- This ensures that findings are likely to be more relevant to those obtained in field trials than anticipated.
 - Relevant transferable protocols for scaling up combined stresses in the large phenotyping platform

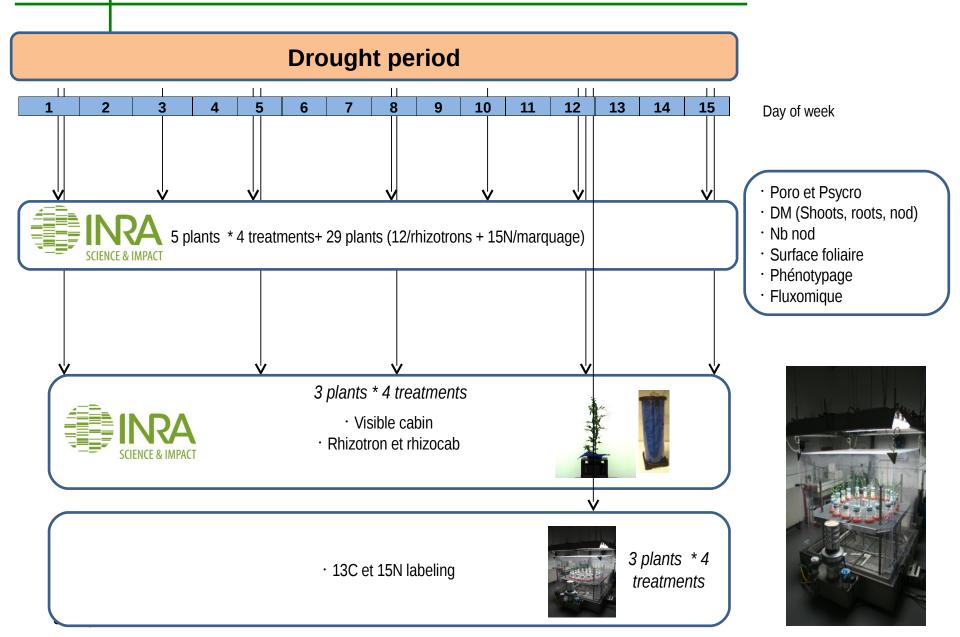
=> Large scaling experiment









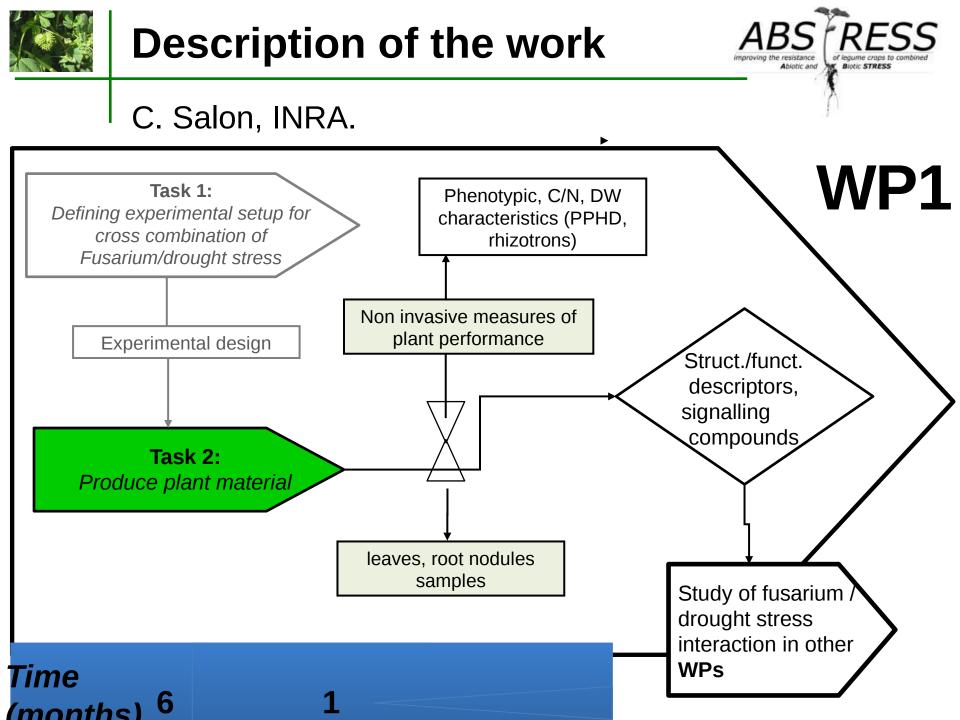




Iasks



efining experimental setup for oss combination of INRA/CSIC <i>Isarium</i> /drought stress
oss combination of INRA/CSIC



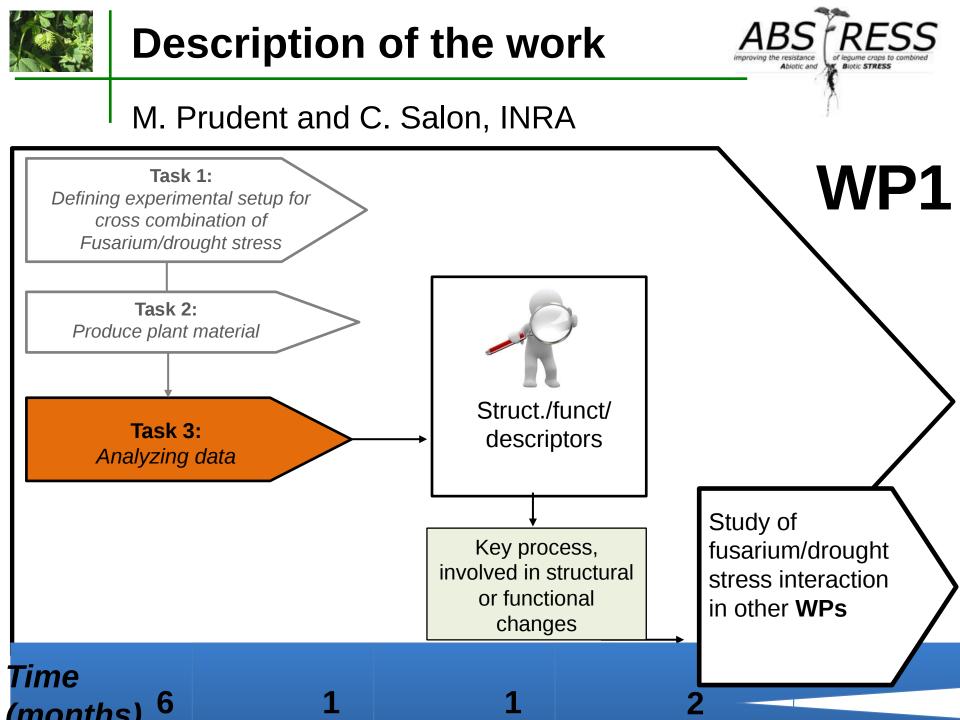


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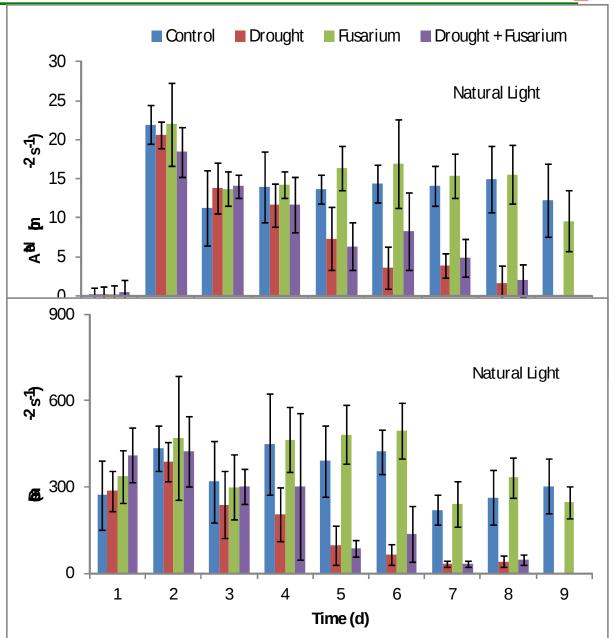


Some results: medicago and

pea

How did treatments affect after 12 days: - development, - growth: carbon acquisition, partitioning &

Physiological data (Pre-pilot)



University of Essex

IRGA measurements (Essex)

Instantaneous (snapshot) readings of Assimilation rate (Anet) and stomatal conductance (gs) under natural light and saturating light. Two leaves

(youngest fully expanded) per plant and five reps per treatment.



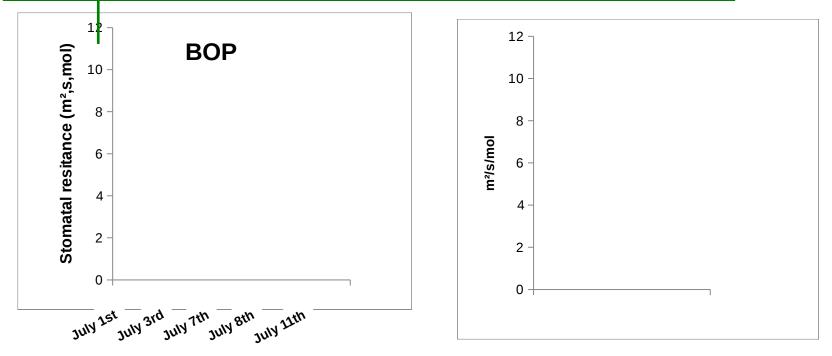
•Example from Pre-pilot.

University of Essex

All IRGA measurements were re-calculated to correct for area. All measurement taken between 9 am -12pm



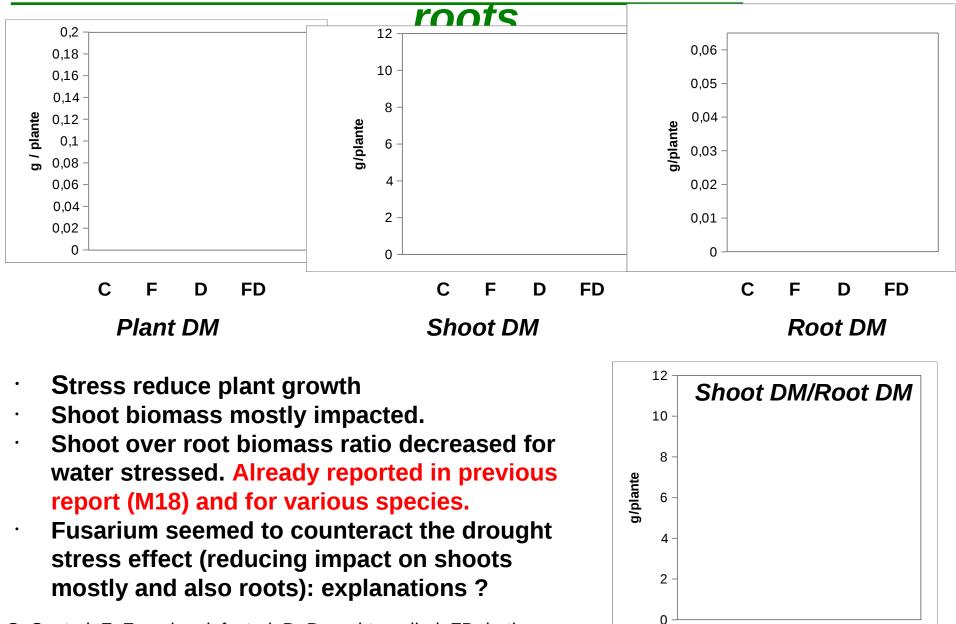
Stomatal resistance



- Drought increases stomatal resistance.
- Fusarium:
 - \cdot amplified the drought stress of plants for both BOM and BOP
 - had no effect when plants were well watered for pea and a detrimental effect similar to water stress alone for Medicago plants



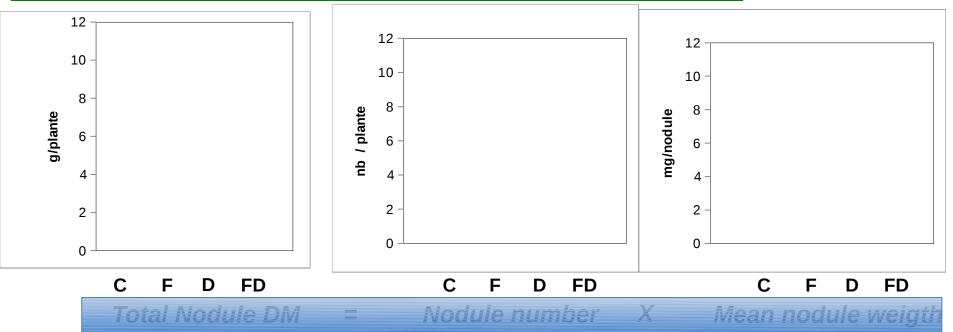
BOMedicago Growth: plant, shoots,



C: Control; F: Fusarium infected; D: Drought applied; FD: both



BOMedicago *Nodules*

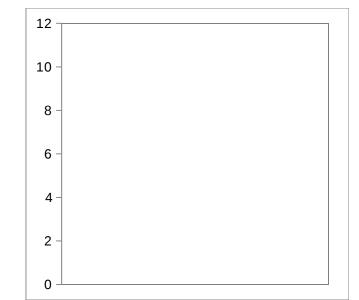


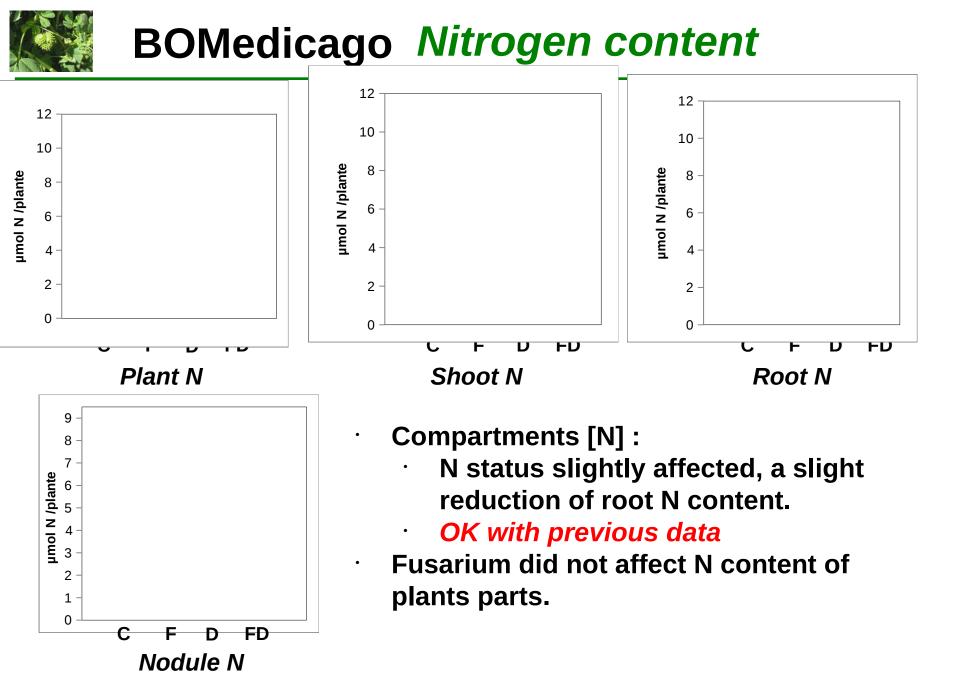
Stress reduce nodule DM (F effect ?)... ...either nodule number and/or ind.

weigth.

- Fusarium leads to bigger nodules in absence of water stress and smaller with drought (?).
- Nodules represent a lower part of nodulated roots under stress

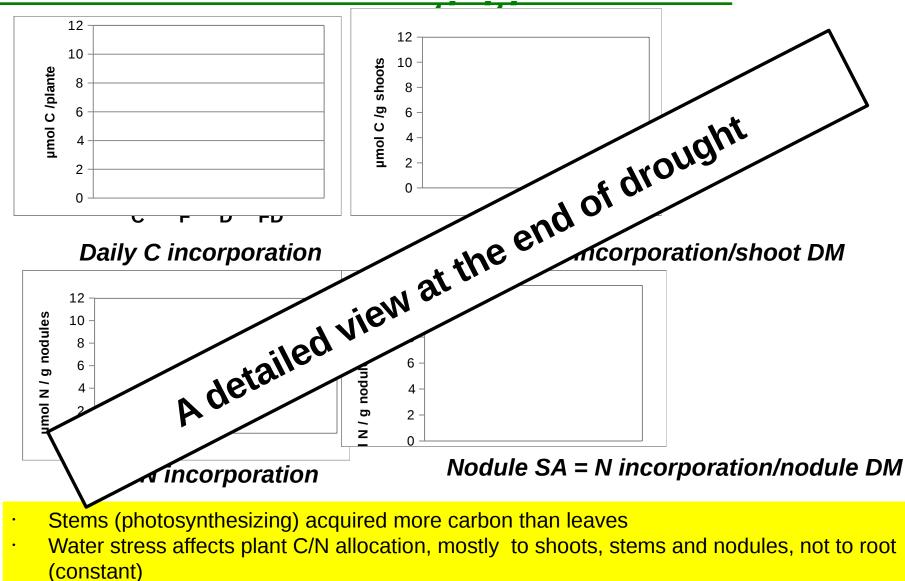
C: Co**Stimilar**uasiun i Mared; D: Drought applied; FD: both





C: Control; F: Fusarium infected; D: Drought applied; FD: both

BOMedicagoFluxo: N and C specific



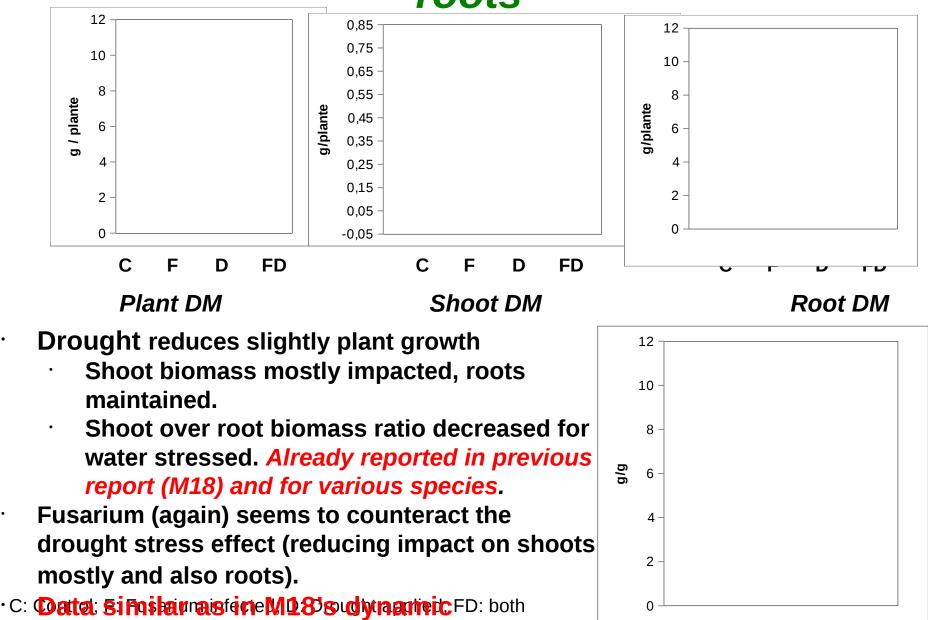
- No effect of Fusarium on C and N acquisition
- Leaf SA reduced, Nodules SA maintained by drought



BOPea

Growth: plant, shoots,

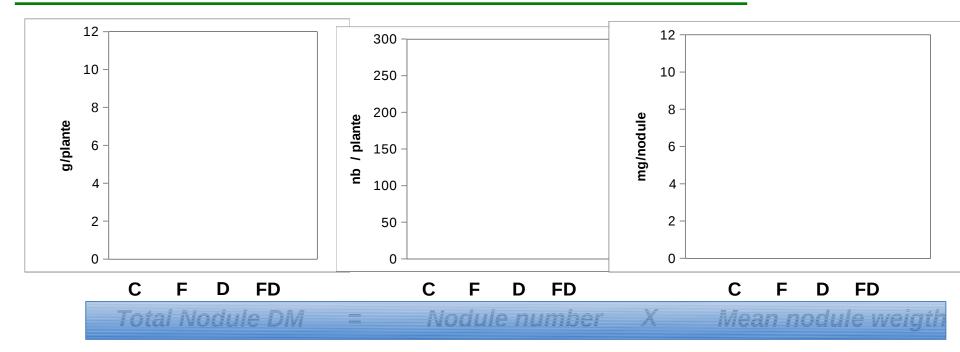








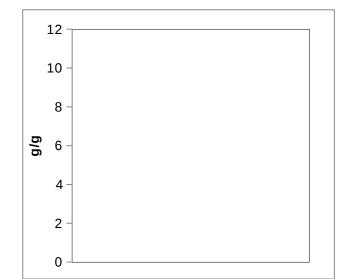
Nodules



Drought

- reduces nodule weigth and number with fusarium.
- Fusarium leads to bigger nodules with or without water stress
- Nodules = a lower part of nodulated roots under drought stress
- Similar as in M18 and for Mt experiment

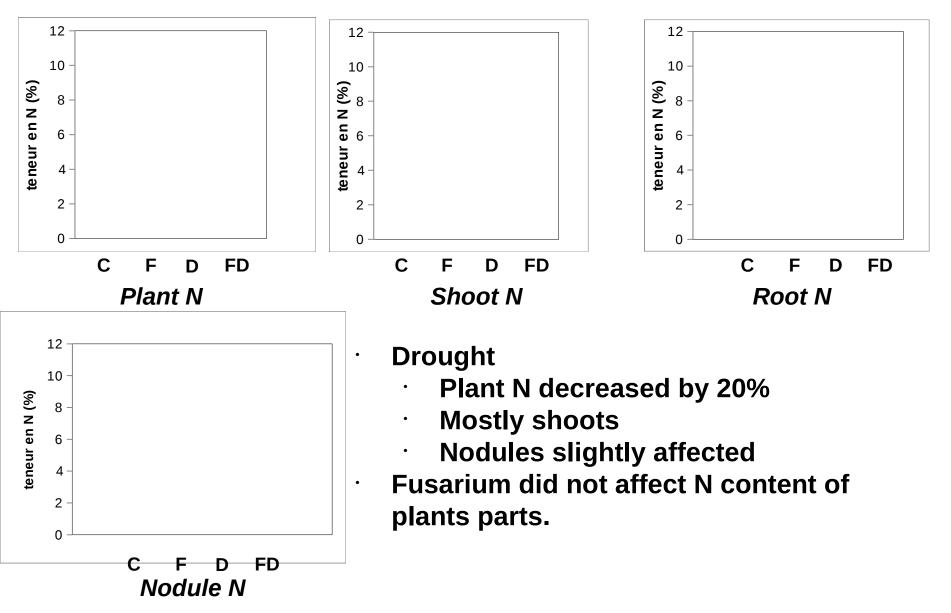
C: Control; F: Fusarium infected; D: Drought applied; FD: both







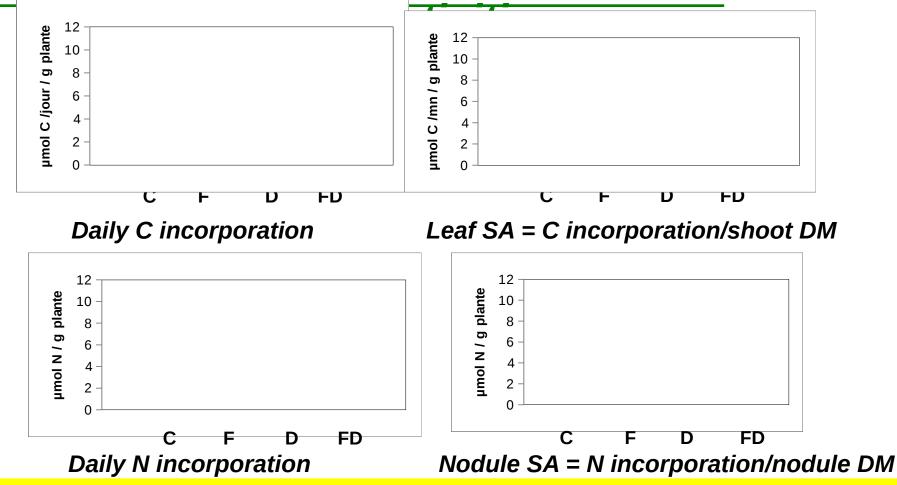
Nitrogen content



C: Control; F: Fusarium infected; D: Drought applied; FD: both



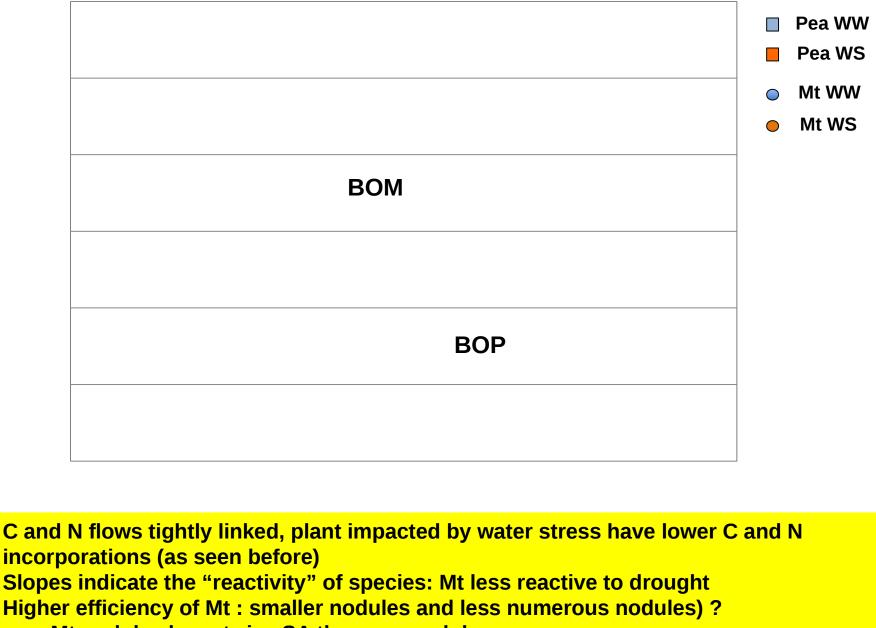
Fluxo: N and C specific



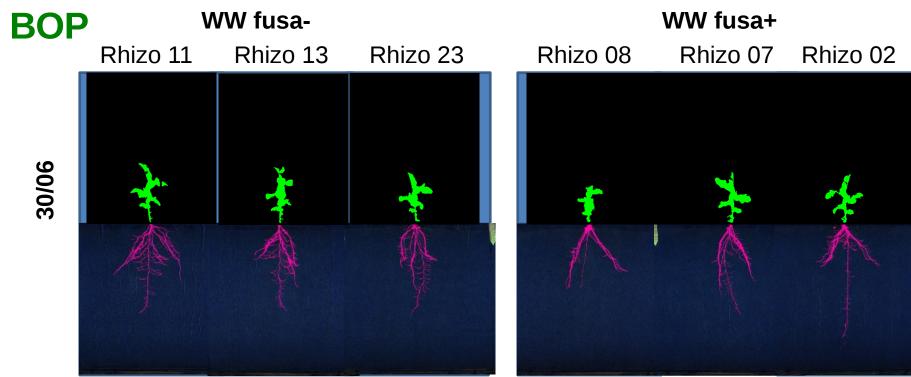
- Pea leaves main sink for new C
- Water stress affects plant C and N allocation, mostly to shoots, stems and nodules, not to root (maintained), *like Mt*
- No effect of Fusarium on C and N acquisition
- Both leaf SA and Nodules SA reduced by drought, reinforced by total nodule biomass decrease As the decrease in nodule biomass was rel. less important than that of total N plant incorporation a greater decrease of nodule specific activity is observed/expected.

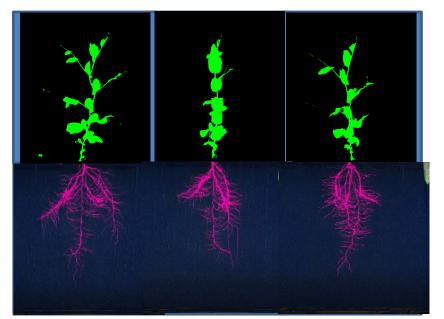


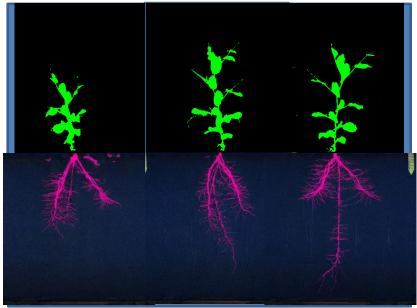
BOPea and BOM



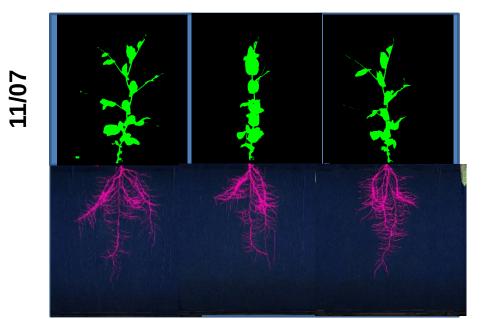
Mt nodules have twice SA than pea nodules

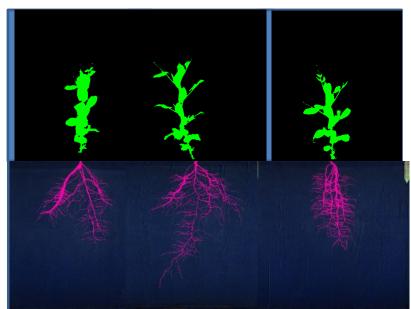






BOP WW fusa-Rhizo 11 Rhizo 13 Rhizo 23 30/06





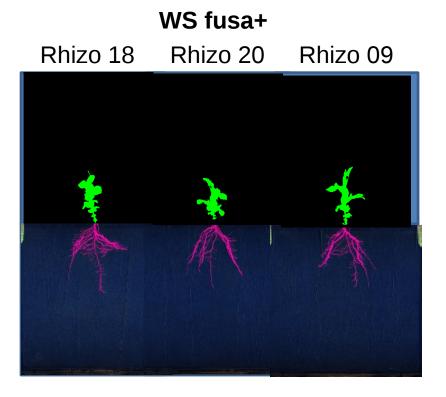
WS fusa-

Rhizo 17

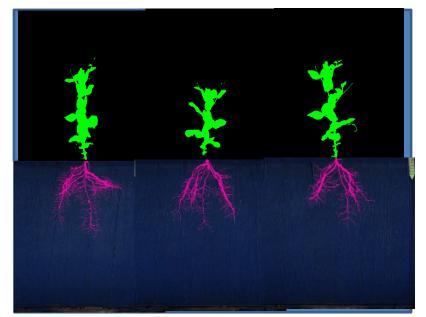
Rhizo 14

Rhizo 24

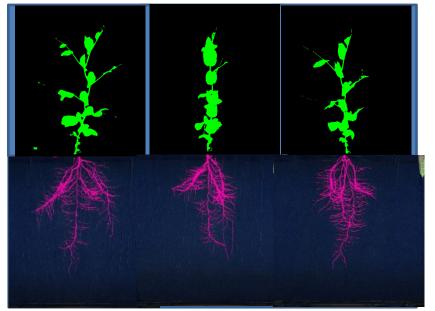
WS fusa-BOP Rhizo 24 Rhizo 17 Rhizo 14 30/06

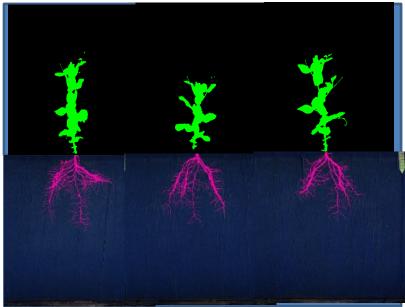


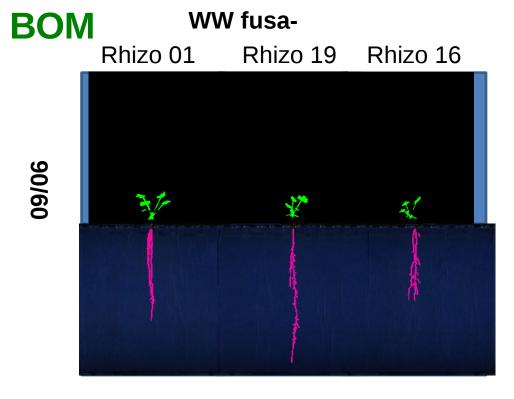


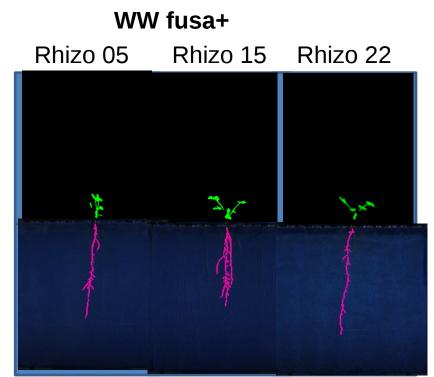


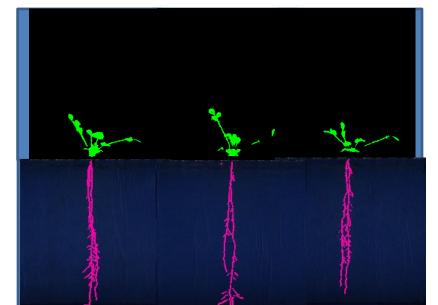
BOP WW fusa WS fusa+ Rhizo 11 Rhizo 13 Rhizo 23 Rhizo 18 Rhizo 20 Rhizo 09 OP Image: Comparison of the state of t

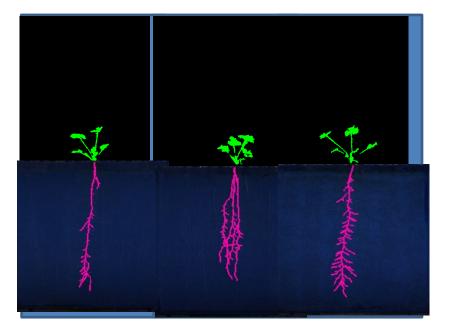


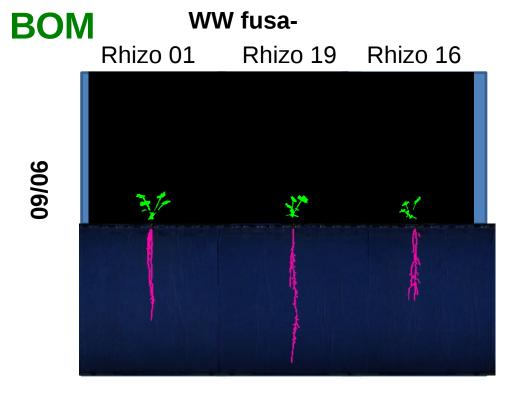


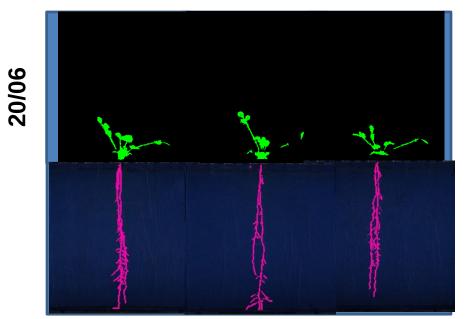


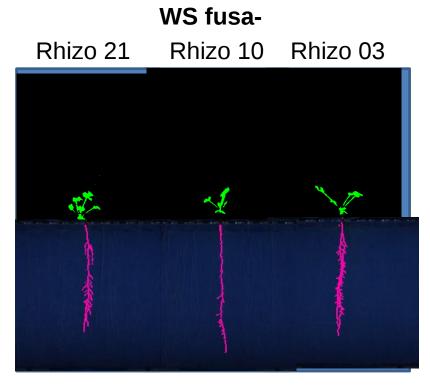


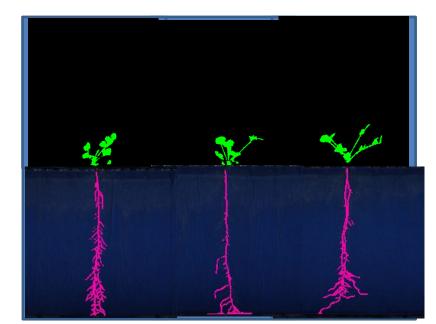


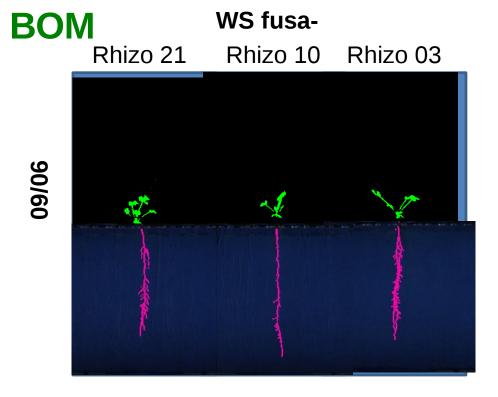


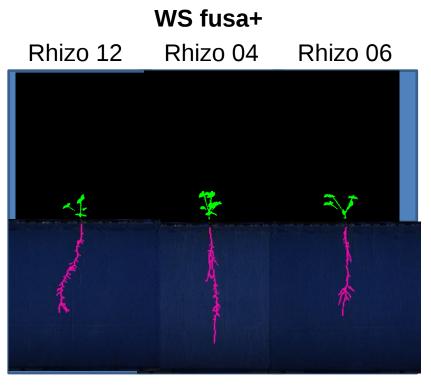


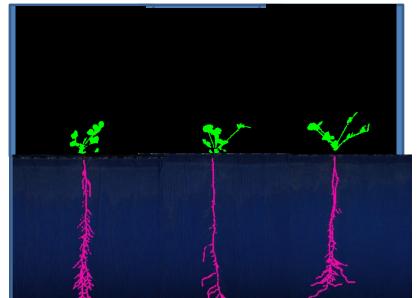


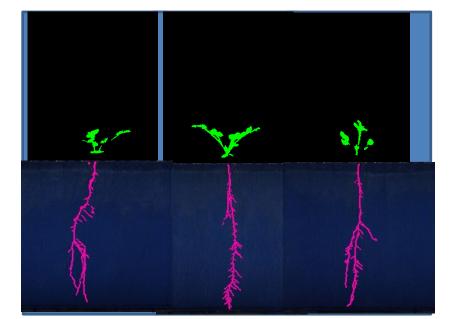




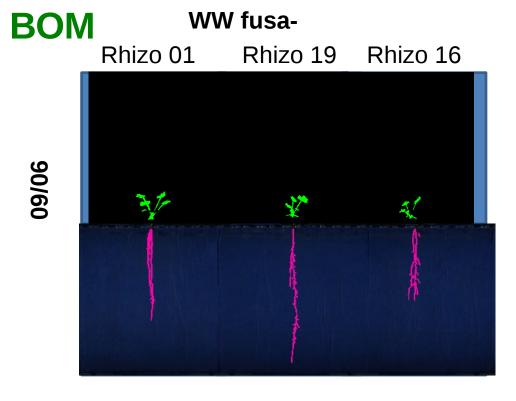


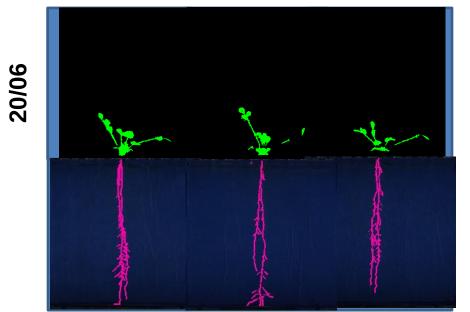


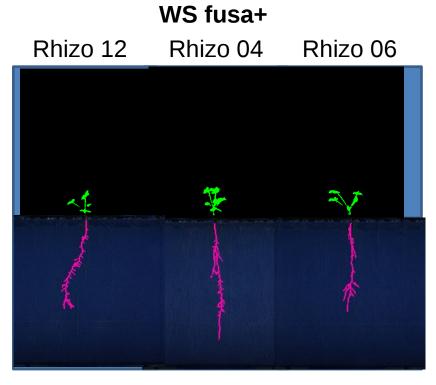


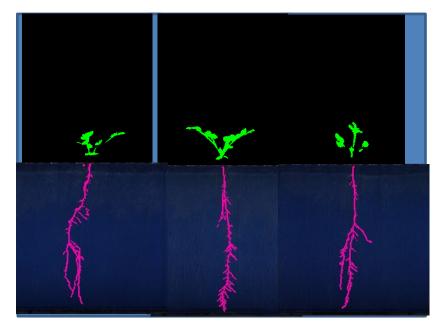


20/06



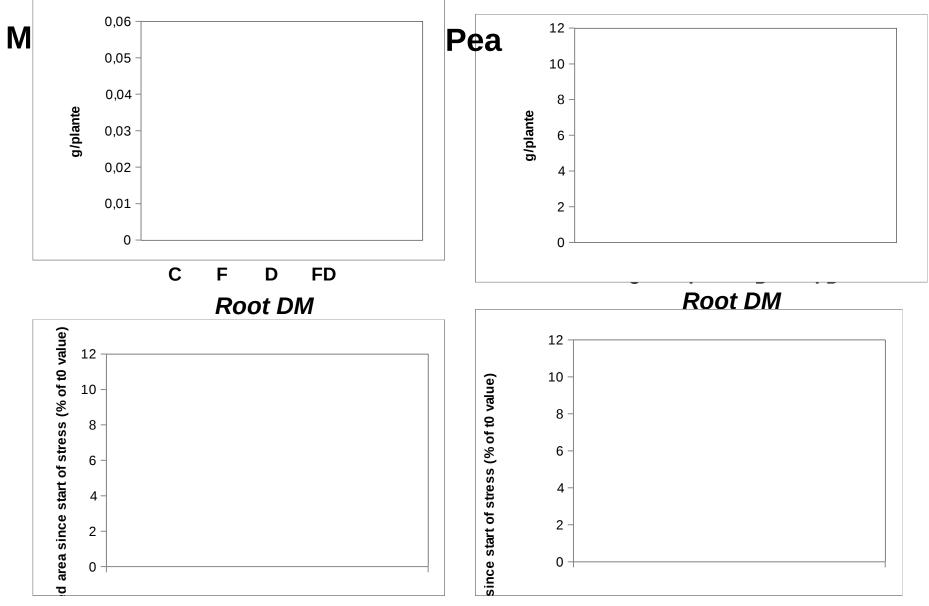








BOM and BOP: Roots RhizoTubes vs



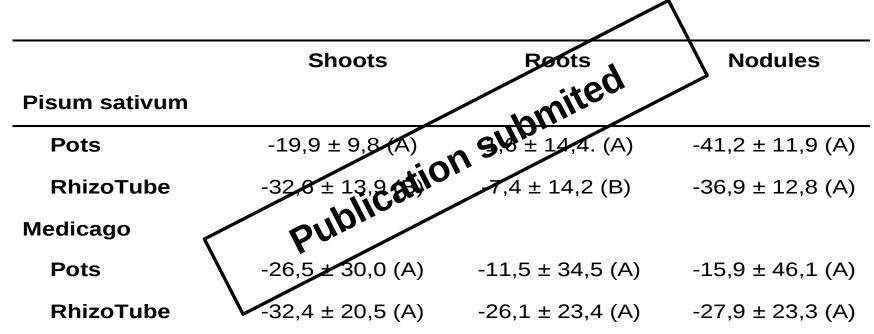
Root Projected area

Root Projected area



BOPea and BOM: RhizoTubes vs

Pots ? Modulation of biomass allocation to shoot, roots and nodules of pea (Kayanne genotype) and Medicago truncatula plants subjected to a water stress versus well irrigated plants. Results are expressed as the ratio of the difference in biomass of shoots (BMS), roots (BMR) or nodules (BMN) between water stress (WS) and well watered (WW) plants to the biomass of well watered plants (n= 5). As an example for shoots, $(BMS_{WS}-BMS_{WW})*100/BMS_{WW}$.



Same results on another experiment with pots or rhiztrons on pea and medicago: RT are usefull tools

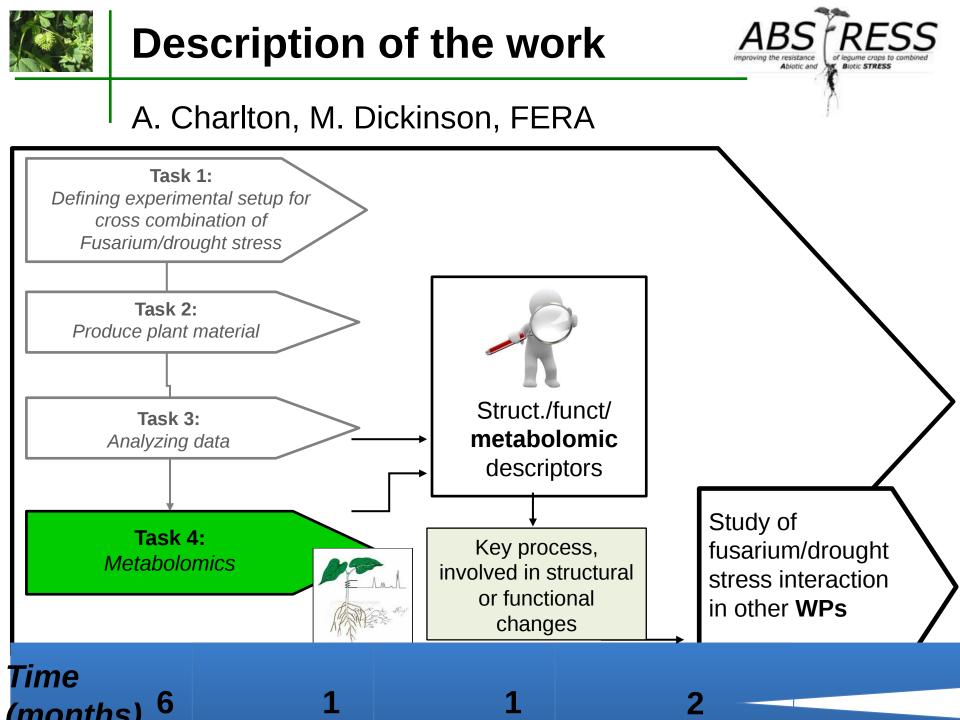


Iasks



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T1.1	Defining experimental setup for cross combination of <i>Fusarium</i> /drought stress	INRA/CSIC	





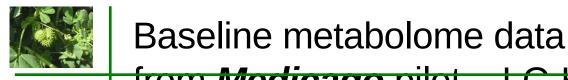


Baseline metabolome data from Medicago

/pea

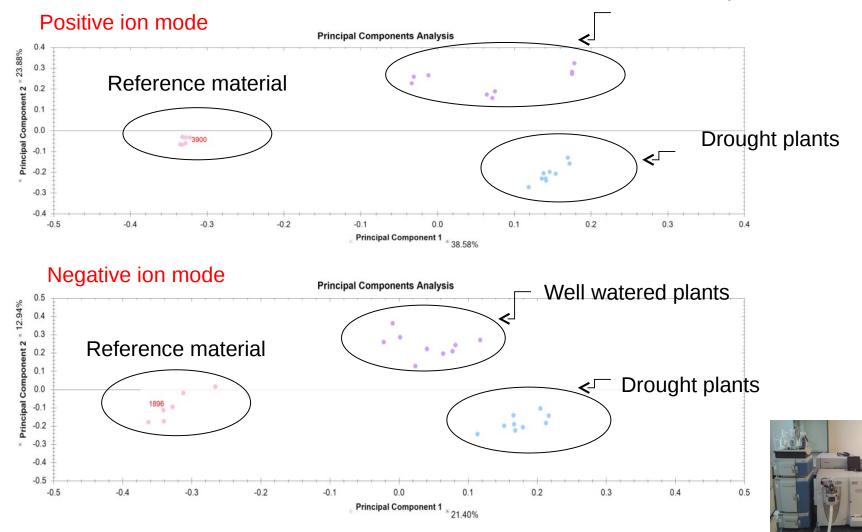






from *Medicago* pilot – LC-HRMS PCA

Well watered plants



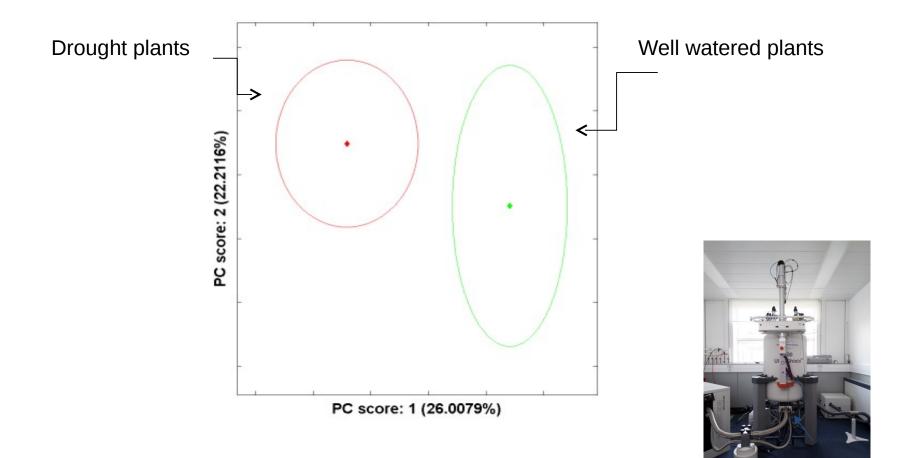


Example compounds signficantly changing in drought stressed *Medicago* tentatively identified by LC-HRMS

			Mean max fold
Tentative compound ID	Levels observed (Up / Down)	lonisation mode	change
putrescine	<u>↑</u>	Positive	68.4
hesperetin (iii)	1	Both	39.1
dihydrobiochanin A (ii)	1	Positive	33.7
glutathione	↓	Positive	31.7
pisatin (i)	1	Positive	31.0
ascorbate	Ļ	Negative	29.4
ferreirin (iii)	1	Both	25.3
4-anisic anhydride (ii)	1	Both	24.9
dehydroascorbate	Ļ	Positive	16.3
ermanin (i)	1	Positive	15.0
proline	↑ (Positive	12.9
cirsiliol	1	Negative	8.2
hematoxylin (iii)	1	Negative	7.8
histidine	1	Positive	6.8
inositol (iv)	Ļ	Positive	6.4
abscisic acid	1	Negative	5.0
raffinose	1	Positive	4.6
glucose-6-phosphate	Ļ	Positive	4.4
hexapyranose (iv)	1	Positive	3.9
oxylipin-3	1	Positive	3.5
malic acid	↓	Negative	3.5
asparagine	1	Positive	3.5
pectolinarigenin (i)	1	Positive	3.4
gamma-tocopherol	1	Positive	2.8



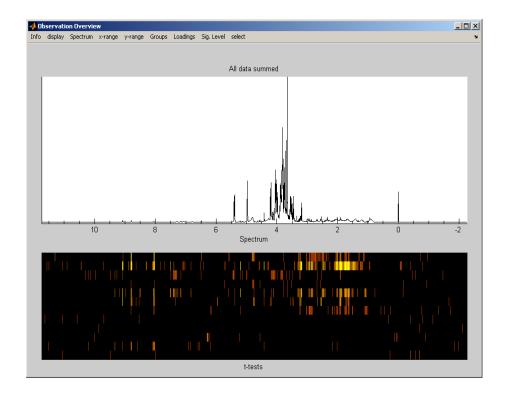
Baseline metabolome data <u>from **Medicago** pilot – LC-HRMS PCA</u>





Metabolite variation in drought stressed vs well watered **pea** study: NMR data

- •Significant t-test results highlighted by brighter colours
- •Clearly metabolic changes associated with drought stress have been detected in the seeds







Metabolite variation in drought stressed vs well watered pea study: NMR data

Metabolite	Chemical shift correlations (ppm)
Proline	4.133-63.87; 3.342-48.72; 3.423-48.68; 2.071-31.55; 2.356-31.61; 2.009-26.41.
Leucine	1.712-42.34; 0.969-24.77; 0.957-23.68.
Isoleucine	1.013-17.24; 0.938-13.55.
Valine	3.615-63.25; 2.289-31.84; 1.049-20.64; 0.999-19.37.
Threonine	4.259-68.56; 1.336-21.90.
γ-aminobutyrate	3.012-41.92; 2.311-36.91; 1.904-26.21.
Homoserine (CID:7799)	3.858-56.13; 3.783-61.40; 2.152-34.82; 2.033-34.78.
Myoinositol (CID:892)	4.061-74.95; 3.624-75.24; 3.544-73.89; 3.270-77.14.
Trigonelline (CID:5570)	9.132-148.40; 8.841-147.50; 8.821-148.57; 8.089-130.17; 4.437-50.92.

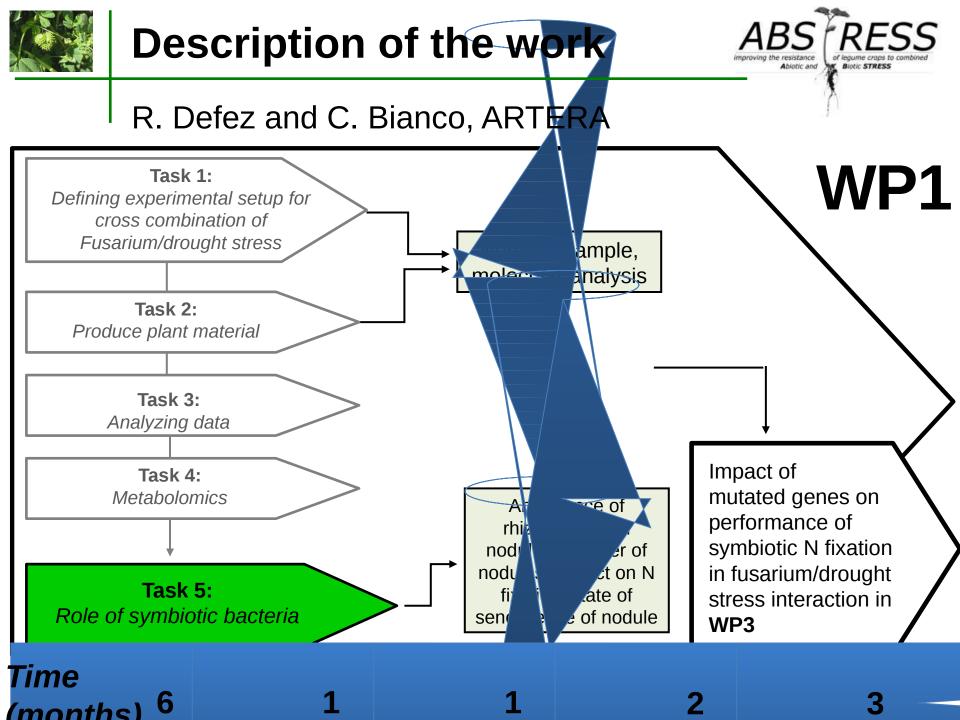


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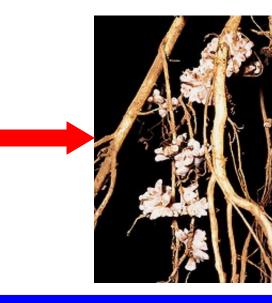


The interaction between legumes and rhizobia leads to the development of aM. truncatulanitrogen-fixing symbiosis



S. meliloti





Root nodules

Sample	Shoot dry wt	Ratio	Seeds wt	Ratio	Pods wt	Ratio
	(mg)		(g/plant)		(g/plant)	

Several environmental factors can adversely affect the performance of symbiotic nitrogen fixation bv legumes. These factors may act at the following levels: survival of rhizobia in the soil, the infection process, nodule growth and nodule function. These factors can also indirectly affect **N2** fixing performance through their negative impact on the host plant growth.

wtr – wild type *rhizobium* GMr – IAA-overproducing *rhizobium* Journal of Experimental Botany, Page 1 of 11 doi:10.1093/jxb/erp140



RESEARCH PAPER

Medicago truncatula improves salt tolerance when nodulated by an indole-3-acetic acid-overproducing *Sinorhizobium meliloti* strain

Carmen Bianco and Roberto Defez*

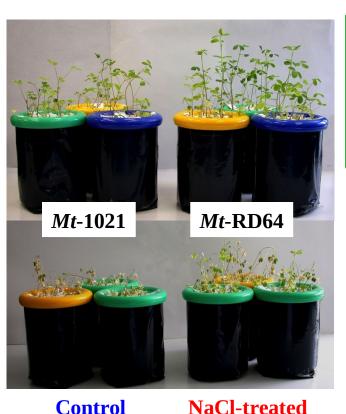
Institute of Genetics and Biophysics 'Adriano Buzzati Traverso', via P. Castellino 111, 80131 Naples, Italy

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(75) In (73) As

Received 12 March 2009; Revised 3 April 2009; Accepted 6 April 2009



Under salt-stress condition, reduced symptoms of senescence, lower expression of ethylene signalling genes, lower reduction of shoot dry weight, and better nitrogenfixing capacity have been observed for Medicago plants nodulated by an IAA-overproducing *S. meliloti* strain.



US007846708B

United States Patent Defez METHOD FOR INCREASING THE SURVIVAL OF BACTERIAL STRAINS OF THE RHIZOBIUM GENUS		(10) Patent No.:(45) Date of Patent:		US 7,846,708 B2 Dec. 7, 2010	
		(52)	C12N 5/02 (2)	006.01) 006.01) 435/243 ; 435/244; 435/252.1; 435/410; 435/420; 435/431	
Inventor:	Roberto Defez, Rome (IT)	(58)	Field of Classification See application file for o	Search None complete search history.	
Assignee:	Consiglo Nazionale delle Richerch, Rome (IT)	(56)	Referenc		
	Kome (11)		U.S. PATENT I	DOCUMENTS	

Vol. 76, No. 14

Improvement of Phosphate Solubilization and Medicago Plant Yield by an Indole-3-Acetic Acid-Overproducing Strain of *Sinorhizobium meliloti*[⊽]†

Carmen Bianco and Roberto Defez* Institute of Genetics and Biophysics "Adriano Buzzati Traverso," Via P. Castellino 111, 80131 Naples, Italy

Received 13 November 2009/Accepted 18 May 2010



R_limitin *Mt*-1021

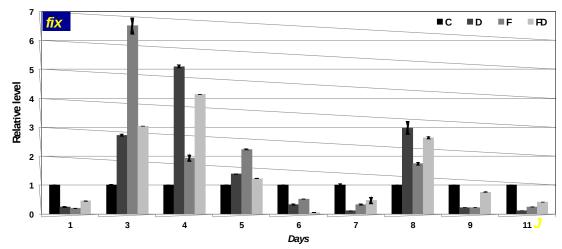
Plimiting Mt-RD64

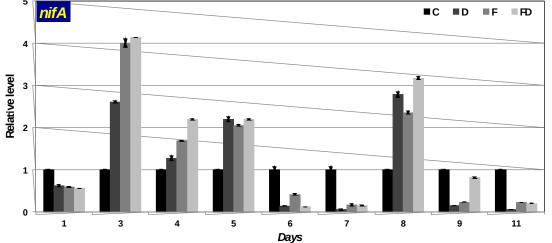
Medicago truncatula plants nodulated by an IAA-overproducing S. meliloti strain and grown under P-deficient conditions showed significant increases in both shoot and root fresh weights when compared to those nodulated by the wild type strain.



(12) United States Patent Defez		(10) Patent No.:(45) Date of Patent:		US 9,157,104 B2 Oct. 13, 2015		
(54)	(4) METHOD TO IMPROVE PHOSPHATE SOLUBILIZATION IN PLANTS		(56)	References Cited		
(75)) Inventor: Roberto Defez, Napoli (IT)		U.S. PATENT DOCUMENTS 7.846,708 B2* 12/2010 Defez 435/24			
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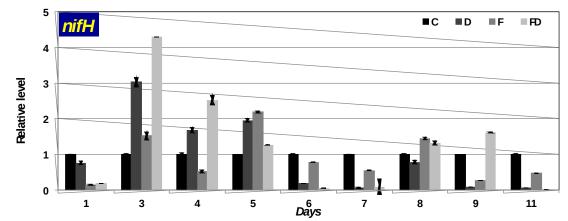
qPCR analysis of nif and fix genes in M. truncatula root nodules during drought stress and Fusarium attack



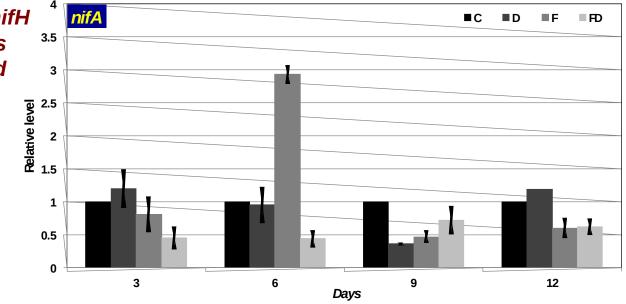


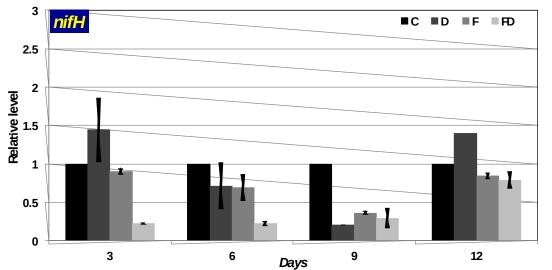
The relative expression levels shown in the Figures are >1 for genes more highly expressed in nodules of D (WS F-), F (WW F+) and FD (WS F+) plants as compared to C (WW F-) control plants. The data reported in the Figure are the means standard deviation of four biological replicates.

The negative effects related to the stress treatments were visible after six days of treatment. Indeed, at this time, all the three genes tested involved in nitrogen fixation were significantly repressed.



qPCR analysis of nifA and nifH genes in Pea root nodules during drought stress and Fusarium attack



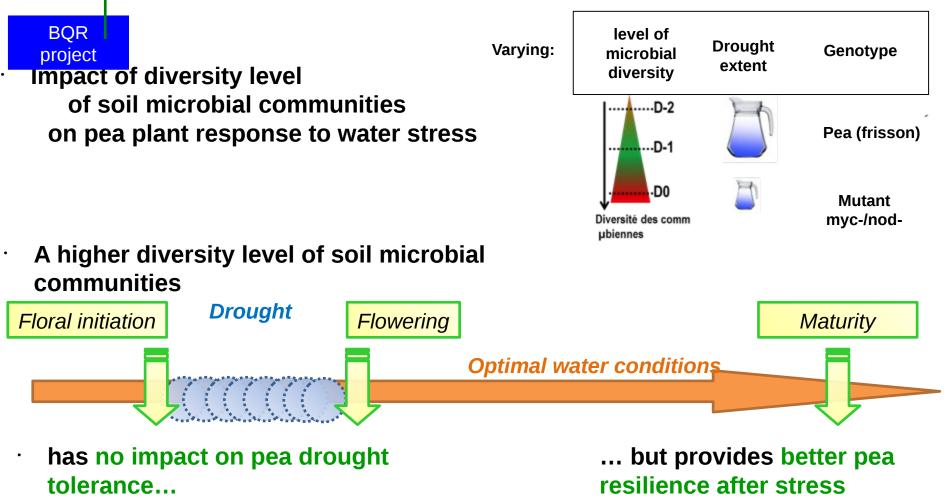


The relative expression levels shown in the Figures are >1 for genes more highly expressed in nodules of D (WS F-), F (WW F+) and FD (WS F+) plants as compared to C (WW F-) control plants. The data reported in the Figure are the means standard deviation of four biological replicates.

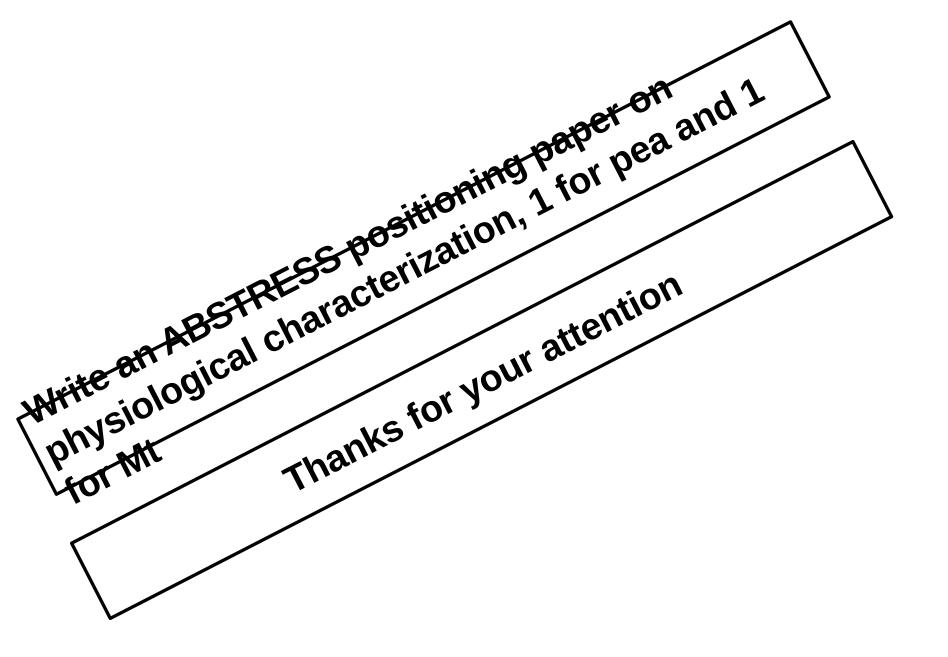
The data clearly show that the singular stress negatively affect the expression of selected genes and that the down-regulation is even more pronounced when double stress (DF) is applied. The down-regulation of N-fixation genes begins as early as the third day of treatment reaching its maximum after 6 day, after which the damage was retained.



Impact of microbes diversity level



Similar response with or without symbioses: non symbiotic communities play a role in this response





Conclusions

- Both integrative and high resolution experiments : drought greatly decreased carbon incorporation of both Medicago and pea, root biomass less impacted.
- Water stress negatively impacted nodule number in Medicago while mean nodule biomass was targeted in pea.
- No clear trend concerning pathogen attack.
- Labelling experiment shown that N flow was greatly reduced by drought for pea (leaves, stems and nodules), while medicago seemed to be much less impacted for its compartments.
- In pea both i) leaf specific activity and ii) nodule biomass and nodule specific activity were severely decreased by water stress, not by fusarium.
- · In Medicago, only leaf specific activity was reduced by water stress.
- A tight carbon/nitrogen relationships was obtained during the labelling experiment:
 - Allows estimating the degree of stress sensed by plants, efficiency to react faced to a stress.

Rhizotubes mimics pot growth

Image analysis is a powerful tool to follow dynamically and automatically, nondestructively shoot and root projected area.