Can we improve the efficiency of biostimulants for nitrogen fertilization by studying their biological effects ?

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In the current situation, agriculture meets demand to feed the growing world population in a changing climate. The most used fertilizers are composed of inorganic nitrogen (N) which is an essential macro-element for plants. For most of them, N is taken up as inorganic N source from the soil (nitrate or ammonium) but their availability can largely vary in soils. However, N production and addition in fields have a strong polluting impact on the environment. There is thus an urgent need for strategies allowing a better N use efficiency (NUE) in crops. The use of biostimulants like protein hydrolysates (PH) is one of them. They have been developed to improve nutrient use efficiency, storage and remobilisation of nutrient elements in crops along with resistance to stresses^{1, 2}. This project aims to better understand how a PH, which contain amino-acids, can improve N fertilization.

All experiments were carried out using Arabidopsis plants grown on vertical plates in vitro with 5mM NO₃⁻ and 1% sucrose. Root growth was measured after 12 days in culture and omics analyses were performed after 14 days in culture. PH is manufactured by the Fertinagro company.



- PH solution (mg/L)

Transcriptomic analysis

NO₃⁻ nutrition Effect of NO_3^- supply: 5mM Stimulating effect of PH at

Number of genes differencially expressed with

Fold change of metabolite concentrations normalized by the mean of all samples

12.5 -



all NO₃-concentrations but stronger effect of PH in nitrogen deficiency

 \rightarrow Next: metabolomic and transcriptomic analysis



- Effect of salt stress: Stimulating effect of PH is maintained in moderate NaCl stress
- \rightarrow Next: characterization of abiotic stress response with PH treatment

ANOVA test (16<n<20 seedlings per condition). Letters indicate a significant difference with the control condition.

PH treatment compared to control condition



1254 up-regulated

1287 down-regulated

Amino acid transporters

role ID name UmamiT46 nodulin MtN21-like transporter family protein AT3G28070 JmamiT47 nodulin MtN21-like transporter family protein JmamiT18 nodulin MtN21-like transporter family protein JmamiT05 nodulin MtN21-like transporter family. AT5G407 transporter aa UmamiT33 nodulin MtN21-like transporter family protein AT4G28040 -0.5 0 0.5 1 Log2(FoldChange)



	ID	name	role	
	AT5G26170	WRKY50	Jasmonic acid pathway	
	AT2G34600	JAZ7	Jasmonic acid pathway	
	AT4G23810	WRKY53	member of WRKY TF	
	AT1G72450	JAZ6	Jasmonic acid pathway	
	AT5G 42650	AOS	Jasmonic acid pathway	
	AT1G75040	PR5	Salicylic acid pathway	
	AT1G19180	JAZ1	Jasmonic acid pathway	
	AT3G25770	AOC2	Jasmonic acid pathway	
	AT2G43590	AT2G43590	PR-3 like gene	
	AT4G19810	CHI-C	Chitinase	
	AT1G32640	ATMYC2	MYC activator	
	AT1G70700	JAZ9	Jasmonic acid pathway	
	At2g03760	ATSOT1	Salicylic acid pathway	
	AT2G43570	AT2G43570	Putative basic chitinase	
	AT3G01970	WRKY45	member of WRKY TF	
	AT5G44420	PDF1.2a	Jasmonic acid pathway	
-1.5 -1 -0.5 0 0.5 Log2(FoldChange)				

- Some amino acid transporters are differentially regulated by PH
- **Down-regulation of NO₃⁻** HATS transporters
- \rightarrow Next: measurement of NO_3^- uptake
- PH influences expression of biotic stress markers
- \rightarrow Next: characterization of biotic stress response with PH treatment

Screening of Arabidopsis mutants hypersensitive or resistant to PH



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