

## Molecular clues on eco-physiological response of lichens to nitrogen

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At Whim, an unmanaged ombrothrophic bog ...



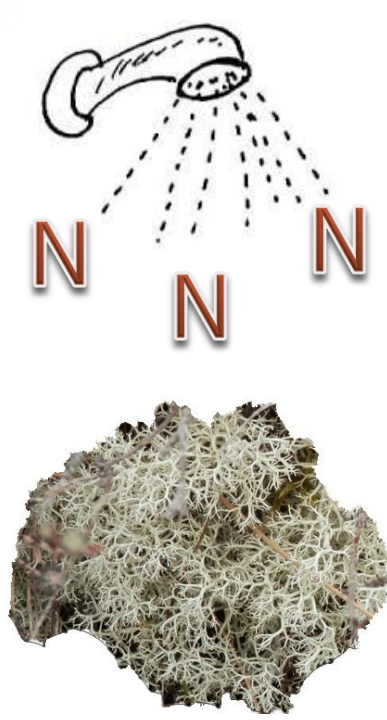
... where N manipulation experiments have been carried on since 2001,...



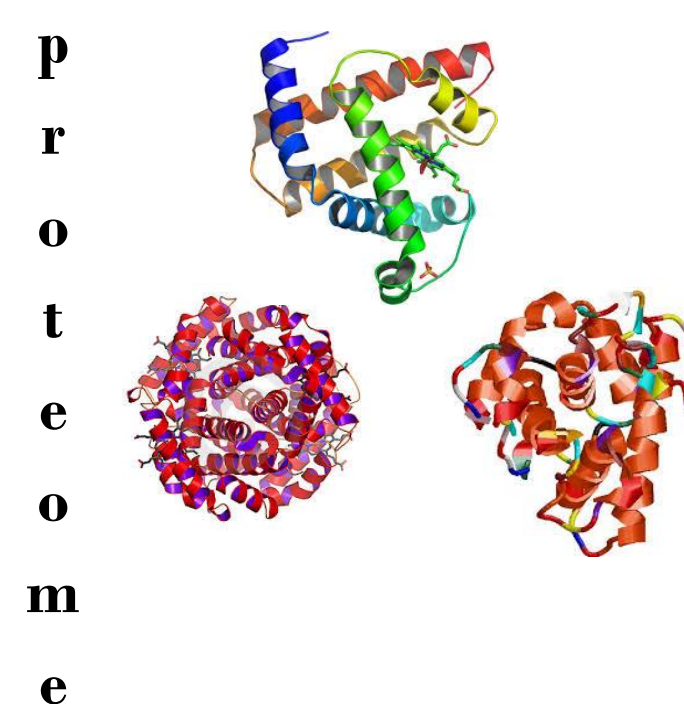
... thalli of *Cladonia portentosa* were transplanted and exposed to nitrogen treatments for 6 months...



... under local climatic conditions.



- 2 forms:  $\text{NH}_4^+$  and  $\text{NO}_3^-$
- 3 doses: 16, 32 and 64 kg N  $\text{ha}^{-1} \text{yr}^{-1}$
- Cumulative exposure of 6 months (short-term) and 11 years (long-term)
- Addition/no addition of phosphorous and potassium: added as  $\text{K}_2\text{HPO}_4$  at a 1:14 PN ratio to give 16 kg N + 0.57 kg P  $\text{ha}^{-1}\text{yr}^{-1}$  and 64 kg N + 4 kg P  $\text{ha}^{-1}\text{yr}^{-1}$

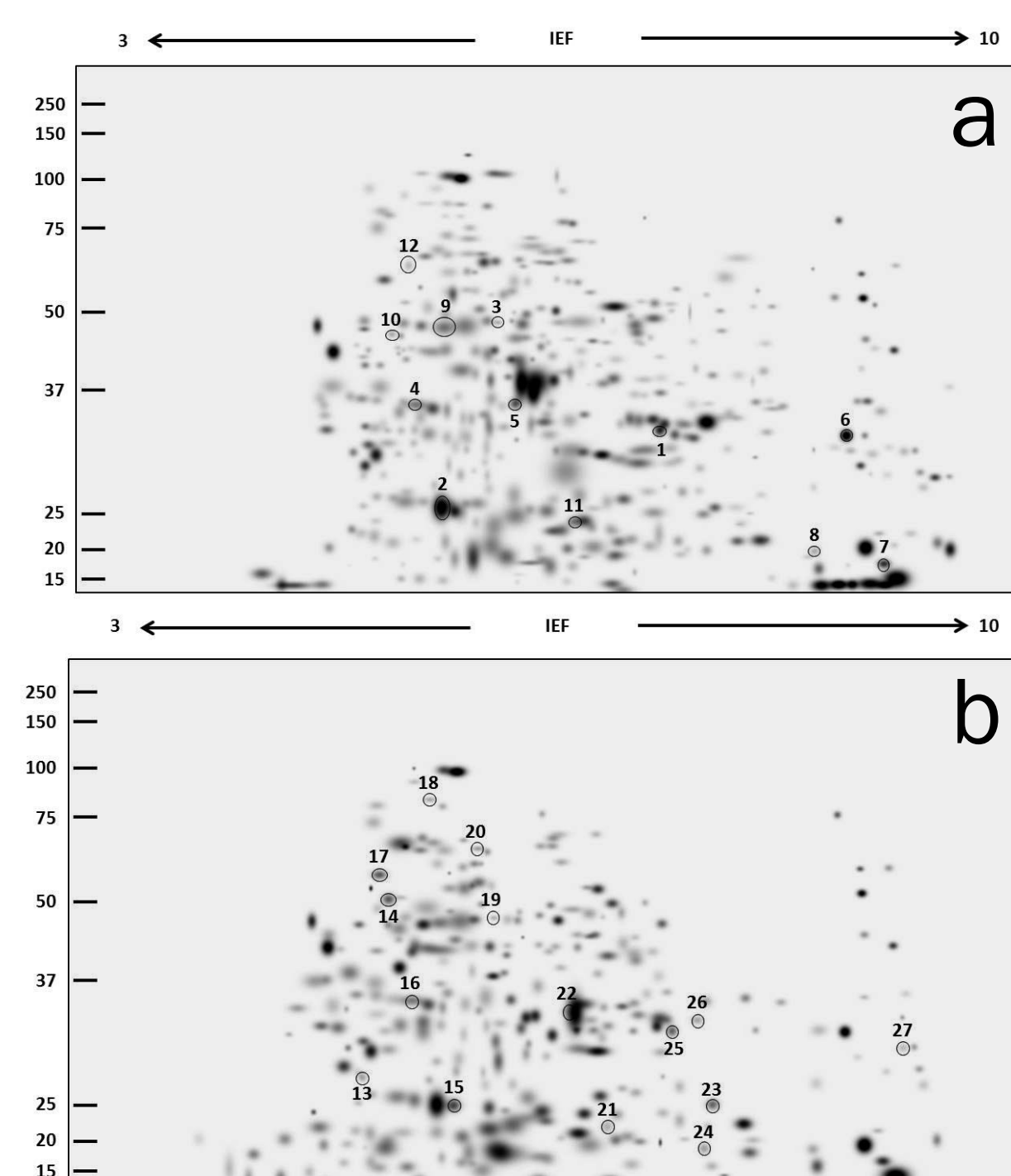


### M&M

The proteome of *C. portentosa* was extracted and analyzed following Nicolardi et al., 2012 (Environmental Pollution 160, 1-10).

Significantly different spots between treatments were selected when comparing: 6-months treatments with different doses of  $\text{NH}_4^+$  and with and without PK; 6-months treatments with different doses of  $\text{NO}_3^-$  and with and without PK.

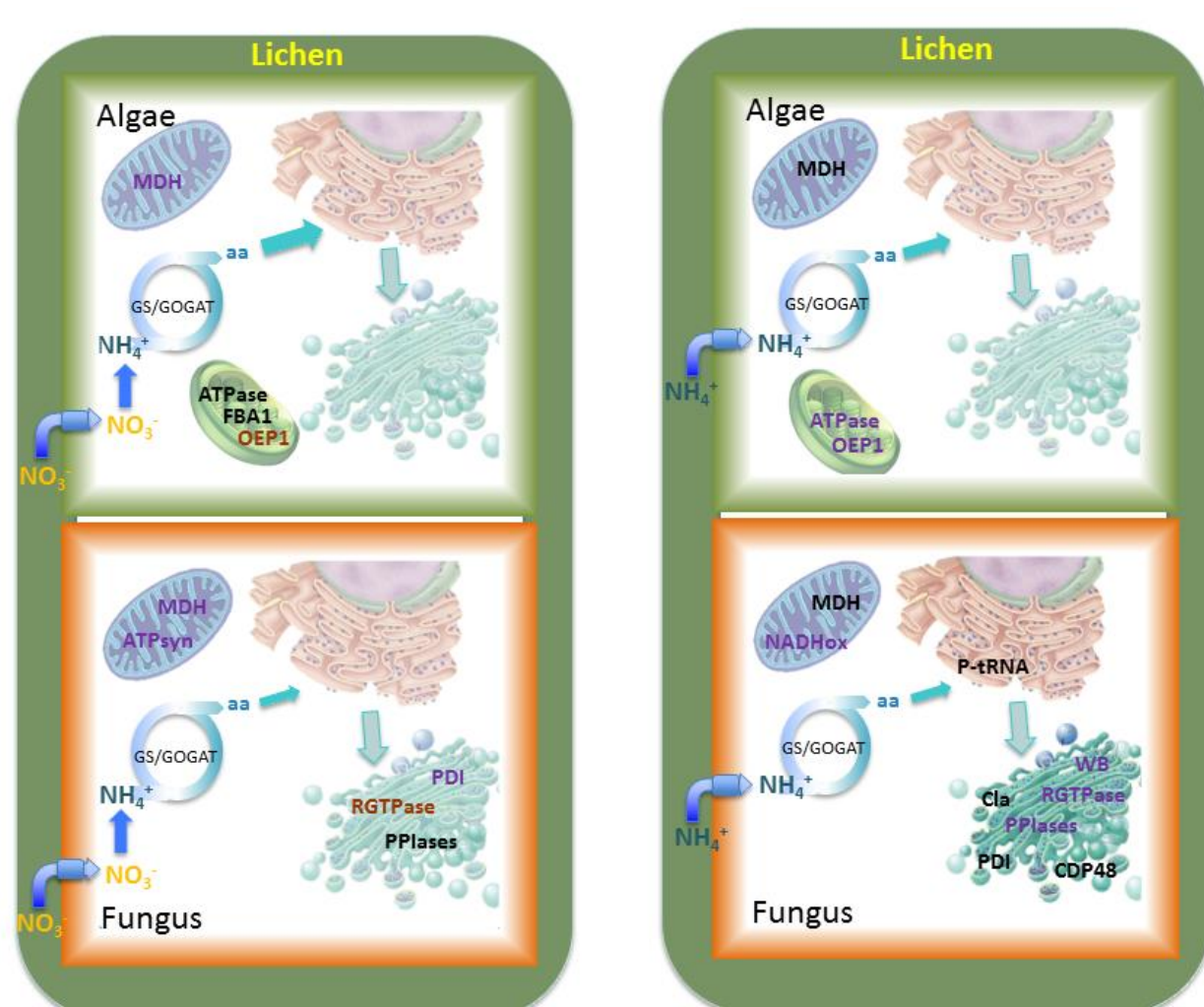
## Results



Master (virtual) gel obtained from the PDQuest-based comparison of 2-D electrophoresis gels of a)  $\text{NO}_3^-$  treated samples and b)  $\text{NH}_4^+$  treated samples. Spots analysed by MALDI-TOF are circled and numbered.

### Summary of changes in protein expression caused by addition of nitrogen

Spot n.	Protein (abbreviation)	Function	Organism	N form	Response to N dose	Threshold	Response to PK
3	ATP synthase subunit beta (ATPase)	Photosynthesis	algae	$\text{NO}_3^-$	upregulated	yes	downregulated
2	Oxygen-evolving enhancer protein 1 (OEPI)	Photosynthesis	algae	$\text{NO}_3^-$	downregulated at 64	yes	downregulated
5	Fructose-bisphosphate aldolase (FBA)	Photosynthesis	algae	$\text{NO}_3^-$	upregulated	yes	downregulated
19	ATP synthase subunit beta (ATPase)	Photosynthesis	algae	$\text{NH}_4^+$	unclear	yes	unclear
15	Oxygen-evolving enhancer protein 1 (OEPI)	Photosynthesis	algae	$\text{NH}_4^+$	unclear	no	downregulated
22	Prolyl-tRNA synthetase	Protein synthesis	bacteria	$\text{NH}_4^+$	upregulated	no	downregulated
12	Protein disulfide-isomerase (PDI)	Folding	fungus	$\text{NO}_3^-$	no	no	upregulated at 64 kg
14-17	Protein disulfide-isomerase (PDI)	Folding	fungus	$\text{NH}_4^+$	upregulated	yes	unclear
24	Peptidyl-prolyl cis-trans isomerase (PPIases)	Folding	fungus	$\text{NH}_4^+$	unclear	no	upregulated at t64 kg
4-10	Ran-specific GTPase-activating protein 1	Regulation	fungus	$\text{NO}_3^-$	downregulated	yes	downregulated at 64 kg
6	Hypothetical protein AN4402.2	Regulation	fungus	$\text{NO}_3^-$	slightly upregulated	yes	downregulated
7	Peptidyl-prolyl cis-trans isomerase (PPIases)	Regulation	fungus	$\text{NO}_3^-$	upregulated	yes	downregulated at 64 kg
27	Putative prohibitin	Regulation	fungus	$\text{NH}_4^+$	downregulated	no	upregulated at 64 kg
13	Clathrin light chain (CLA)	Secretion	fungus	$\text{NH}_4^+$	upregulated	no	downregulated
16	Ran-specific GTPase-activating protein 1	Secretion	fungus	$\text{NH}_4^+$	unclear	yes	unclear
20	Outer membrane autotransporter putative	Secretion	amoeba	$\text{NH}_4^+$	unclear	yes	downregulated
18	Cell division control protein 48 (CDP48)	Protein degradation	fungus	$\text{NH}_4^+$	upregulated	yes	unclear
23	Woronin body major protein (WB)	Stress response	fungus	$\text{NH}_4^+$	unclear	unclear	unclear
11	Thiol-specific antioxidant protein 3	Stress response	fungus	$\text{NO}_3^-$	no	no	upregulated
8	Hypothetical protein DSC_01120	Stress response	bacteria	$\text{NO}_3^-$	upregulated	yes	downregulated
1	Malate dehydrogenase (MDH)	Respiration	algae, fungus	$\text{NO}_3^-$	unclear	unclear	unclear
9	ATP synthase subunit beta (ATPase)	Respiration	fungus	$\text{NO}_3^-$	no	yes	downregulated at 64 kg
25-26	Malate dehydrogenase (MDH)	Respiration	algae, fungus	$\text{NH}_4^+$	upregulated	yes	upregulated at 64 kg
21	NADH-quinone oxidoreductase (NADHox)	Respiration	fungus	$\text{NH}_4^+$	unclear	unclear	unclear



Summary of changes in protein expression (black=upregulation, brown=downregulation, violet=no defined pattern). Abbreviations: aa=aminoacids, GS/GOGAT=glutamate synthase/glutamine oxoglutarate aminotransferase; see table for proteins.

### MOLECULAR RESPONSE

The photobiont showed upregulation of energetic metabolism and photosynthesis

The majority of the under/over expressed proteins belong to the fungus and are involved in protein transportation and regulation (specific tolerance mechanisms?)

Several protein showed a bell-shaped curve with highest expression at the intermediate concentration (existence of a threshold for N)

Upregulation of enzymes involved in protein degradation in case of  $\text{NH}_4^+$  but not  $\text{NO}_3^-$  (futile cycle like in higher plants?)

### ECO-PHYSIOLOGICAL OBSERVATIONS

Analysis of chlorophyll *a* fluorescence parameters showed no impairment of algal functionality

Lichen species with the same photobiont can have different tolerance degrees to N

Fertilizing effect of a small amount of traffic related N

Repeated exposures can exacerbate the toxic effects of N

$\text{NH}_4^+$  is more harmful to lichens than  $\text{NO}_3^-$

