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RESPONSES TO PHOSPHATE DEPRIVATION IN WHITE CLOVER (TRIFOLIUM REPENS L.)

A thesis presented in partial fulfilment of the requirements for the degree of

Master of Philosophy in Plant Biology

at Massey University Palmerston North New Zealand

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2007

Abstract

Four breeding lines of white clover (Trifolium repens L.) were obtained from AgResearch Grasslands, Palmerston North, New Zealand, that had been shown previously to differ in terms of specific growth responses to added phosphate (P) in the field. These were designated Breeding Line (BL) 43 (low performer on low P; low performer on high P), BL 45 (low performer on low P; high performer on high P), BL 47 (high performer on low P; high performer on high P), and BL 49 (high performer on low P; low performer on high P). These breeding lines and five selected genotypes that were propagated from each line (designated 43-7, 43-8, 45-14, 45-4 and 47-9) were rooted in half-strength Hoagland solution in vermiculite for two weeks and then transferred to half-strength Hoagland liquid media for five weeks prior to the initiation of the experiments. For the breeding line screening, plants were acclimatized in a constant temperature environment for one week prior to treatments, while for the genotypic screening, plants were maintained in a temperature-controlled glasshouse. These lines and genotypes were characterized in relation to P uptake and utilization efficiency by growing in P-sufficient media (+P; 0.5 mM KH₂PO₄) and P-deficient media (-P; 0 mM KH₂PO₄) for 3, 5, 7 and 14 days (for the breeding line screening) and 7, 14 and 21 days (for the genotype screening). Over the time course, inorganic phosphate (Pi) content in leaves, non-specific acid phosphatase (APase) activity in intact roots (both as a total soluble activity and a cell-wall-associated activity), isoenzyme analyses, shoot dry weight (DW) and fresh weight (FW), leaf area, weight of an individual leaf (designated as the weight of the first fully expanded leaf), root FW, and the root:shoot (R:S) ratio were determined.

Pi deprivation enhanced the induction of one major low mobility cell wall acidic isoform, two minor high mobility cell wall acidic isoforms and one major low mobility cell wall basic isoform in all genotypes. Furthermore, the activity of one major low mobility cell wall basic isoform was more higher in genotype 45-14 and one minor high mobility cell wall basic isoform was induced only in genotype 45-14 in response to Pi deprivation.

In terms of individual BLs and genotypes, the screening results showed that BL 49 and genotype 45-14 displayed a constant Pi content and a slow induction of APase activity in the –P media, and had the highest total biomass FW in both +P and –P media.

Overall (in both treatments) BL 49 and genotype 45-14 are the most efficient at utilizing available P as they produced the largest biomass FW, produced more roots in P-deprived media when compared with the other BLs and genotypes, and were more efficient in utilizing the P for the synthesis of biomass. BLs 43 and 45 and genotypes 43-7 and 43-8 are less efficient at utilizing available P, while under P deprivation, BL 45 and genotype 45-14 are the most efficient at utilizing P compared to the other BLs and genotypes. The study also showed that the Pi content in leaves and APase activity in roots was found to be the plant parameter most sensitive to Pi deprivation, and the results suggest that the selection of white clover germplasm for satisfactory performance under low P availability can be carried out using these two parameters as criteria.

Acknowledgments

I would like to thank my senior supervisor, Prof. Michael T. McManus, for his excellent supervision, patience, and encouragement throughout my study and during the preparation of this thesis.

I would like to thank my co-supervisor, Dr. Derek Woodfield, for sharing his knowledge and expertise in white clover, in addition to his guidance, advise, discussion and construction comments.

Other very important contributions were made by the member of the lab, both past and present. They are Dr. Balance Chen, Dr. Ning, Dr. Sarah Dorling, Jan, Susanna, Ludivine, Marissa, Aluh, Elizabeth, Rachel, Fiona, Mathew, Matthew, Lena, and Cait.

I wish to thank all of my friends for their wonderful companionship and support throughout my study at Massey University.

I wish to thank Rosemary Knowles and Mark Herrings for their wonderful friendship during the first two years I was in New Zealand.

I wish to express my sincere gratitude to the NZAID for funding my study in New Zealand, and Pattimura University (UNPATTI) Ambon, Indonesia for granting permission to study here. Similar appreciations go to the Institute of Molecular Biosciences and New Zealand Society of Plant Physiologists for providing generous travel and accommodation assistance grant to make it possible for me to go to a conference in Christchurch and in Australia.

Finally, I would like to thank my parents and Ulf who always support me through their love, encouragement and understanding.

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Abbreviations

 $A_{405 \text{ nm}}$ absorbance $[\log(Io/I)]$ in a 1 cm light path at 405 nm

3-PGA 3-phosphoglyceric acid

APase acid phosphatase

APS ammonium persulphate

AtACP5 Arabidopsis thaliana acid phosphatase type 5

AtPAP12 Arabidopsis thaliana purple acid phosphatase type 12

AVG aminoetoxyvinylglycine

BL breeding line

BM biomass

cDNA complementary deoxyribonucleic acid

cv cultivar d day

DNA deoxyribonucleic acid

DOT days of treatment

DTT dithiothreitol

DW dry weight

FW fresh weight

G genotype

g gram

g acceleration due to gravity (9.81 m s⁻²)

h hour

kD kiloDalton (unit of molecular mass)

kPa kiloPascal

L litre

Lycopersicon esculentum phosphate starvation-induced

gene type 2

M molar, moles per litre

mg milligram

MilliQ water water that has been purified by passing through a MilliQ

ion exchange column

 $\begin{array}{ccc} \mu g & microgram \\ \mu M & micromolar \\ miR & microRNA \\ mL & milliliter \\ mm & millimeter \\ mM & millimolar \end{array}$

NIL near isogenic line

nm nanometer

NS not significant °C degree Celsius

OD optical density at x nm in a 1 cm light path

PAE phosphorus acquisition efficiency
PAGE polyacrylamide gel electrophoresis

PEP phospho(enol) pyruvate

PGA phosphoglycerate

Pht phosphate transporter
Pi inorganic phosphate

psr1 phosphate starvation response type 1
PUE phosphorus utilization efficiency

pup1 phosphate under-producer mutant type 1

R:S root per shoot

RNA ribonucleic acid

RNase ribonuclease

RO reverse osmosis

s second

S-APase secreted acid phosphatase

SPT2 Saccharomyces cerevisiae phosphate transporter type 2

TEMED N,N,N',N'-Tetramethylethylenediamine

Tris tris (hydroxymethyl) methylamine

V Volt (kg m² s⁻³ A⁻¹)

v/v volume per volume

W Watt (kg m² s⁻³)

w/v weight per volume

ρNPP ρ-nitrophenyl phosphate