## **Supplementary material**

Table S1. ANOVA summary table for all response variables.

	P	C	P*C
Colonisation (pot A)	ns	***	ns
Colonisation (pot B)	**	***	*
SDW	***	ns	**
RDW (total)	***	ns	*
Root length (total)	***	**	*
Shoot P content	***	ns	***
Root P content (total)	***	***	***
Root P conc. (pot A)	***	***	***
Root P conc. (pot B)	***	***	***
Pot A P uptake	**	**	***
Shoot PO <sub>4</sub> conc.	***	**	ns

(B)

	G	С	$G^*C$
Colonisation (pot A)	ns	***	ns
Colonisation (pot B)	***	***	***
SDW	***	ns	*
RDW (total)	**	ns	ns
Root length (total)	ns	**	ns
Shoot P content	***	ns	***
Root P content (total)	***	**	***
Root P conc. (pot A)	***	***	***
Root P conc. (pot B)	***	***	***
Pot A P uptake	**	***	***
Shoot PO <sub>4</sub> conc.	*	ns	*

Factors in the analysis were (A) P and C, (B) G and C. Both main effects and interaction terms are indicated. P Soil P Addition, C Pot Configuration, G Genotype. P<0.05; P<0.05; P<0.01; P<0.001.

Table S2. Mycorrhizal colonisation (% of root length) of *mtpt4* mutant genotype, grown at 20 mg P kg<sup>-1</sup>. Values are mean  $\pm$  standard error, n=5. Means followed by the same letter were not significantly different at the P<0.05 level (Tukey's HSD).

		M,M	NM,M	NM,NM
Pot A	mean	35.6	0	0
	s.e.	4.9	0	0
Pot B	mean	35.2	31.7	0
	s.e.	5.9	3.7	0

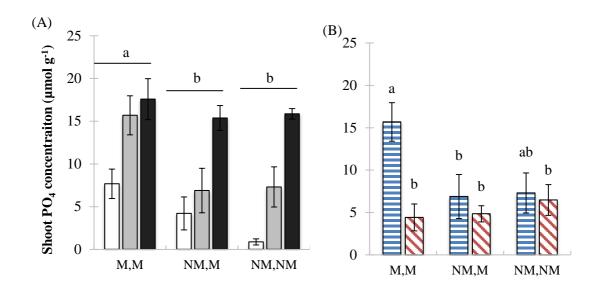


Figure S1. Inorganic phosphate (PO<sub>4</sub>) concentration in shoots.

PO<sub>4</sub> concentration ( $\mu$ mol g<sup>-1</sup>) in the fresh shoots of (A) wild-type plants at three soil P additions: 0 (white bars), 20 (grey bars) and 50 (black bars) mg P kg<sup>-1</sup>, and in (B) wild-type (horizontal bars) and *mtpt4* mutant (diagonal bars) at P20 only. Values are mean  $\pm$  standard error, n=4. Means followed by the same letter were not significantly different at the P<0.05 level (Tukey's HSD). See text for details of statistical analysis.