

Supplemental Table 5. Results of three-way ANOVAs on *NtPT4* and *NtPT1* transcript levels with the categorical factors tobacco lines (four levels: WT, W6:CKX1, 35S:CKX1 or 35S:CKX2), *R. irregularis* RI (two levels: -, +), and *R. irregularis* FM (two levels: -, +). For the mean values see Fig. 5.

Factors	df	<i>NtPT4</i>		<i>NtPT1</i>	
		<i>F</i>	<i>P</i>	<i>F</i>	<i>P</i>
Tobacco line (T)	3	11.9	***	1.2	
<i>R. irregularis</i> RI	1	22.4	***	7.9	**
<i>R. irregularis</i> FM	1	55.3	***	2.6	
T x RI	3	1.9		2.2	
T x FM	3	4.0	*	4.4	*
RI x FM	1	43.6	***	0.0	
T x RI x FM	3	1.2		4.2	*
Residuals	30				

NtPT4, *Nicotiana tabacum phosphate transporter4*. *NtPT1*, *Nicotiana tabacum phosphate transporter1*. WT, wild type. df, degrees of freedom.

For $P < 0.05$, 0.01 and 0.001, significance levels of *F* values are presented as *, ** and ***, respectively, and are in bold. *F* values accompanied by (*) are marginally non-significant and are in italic.