Table S2. Statistics of the reads for each condition mapped onto the reference genome of *P. vulgaris* using the Burrows-Wheeler Aligner (BWA).

	Control	*PvRbohB-*RNAi	Control_Rhiz*	*PvRbohB-*RNAi_Rhiz	Control_Myc*	*PvRbohB-*RNAi_Myc
Percentage of mapped reads	98.31–98.83	97.82–98.68	96.48–98.18	95.85–96.85	97.09–98.55	98.58–98.90
Total number of mapped genes	23,135	23,100	22,393	22,410	22,559	23,357

*Rhiz, inoculated with *R. tropici*; Myc, inoculated with *R. irregularis*.