Figure S1 Intrinsic variability within the data groups. Multidimensional Scaling plots (MDS) (a-c) and Pearson correlation (d-f) of the different replicas per condition of control and PvRbohB-RNAi samples of the transcriptome uninoculated (a, d), inoculated with rhizobia (b, e), and inoculated with AM (c, f). Ctrl: control roots; Bi: PvRbohB-RNAi; Rhiz: inoculated with R. tropici; Myc: inoculated with R. irregularis; 1: replicate 1; 2: replicate 2; 3: replicate 3.