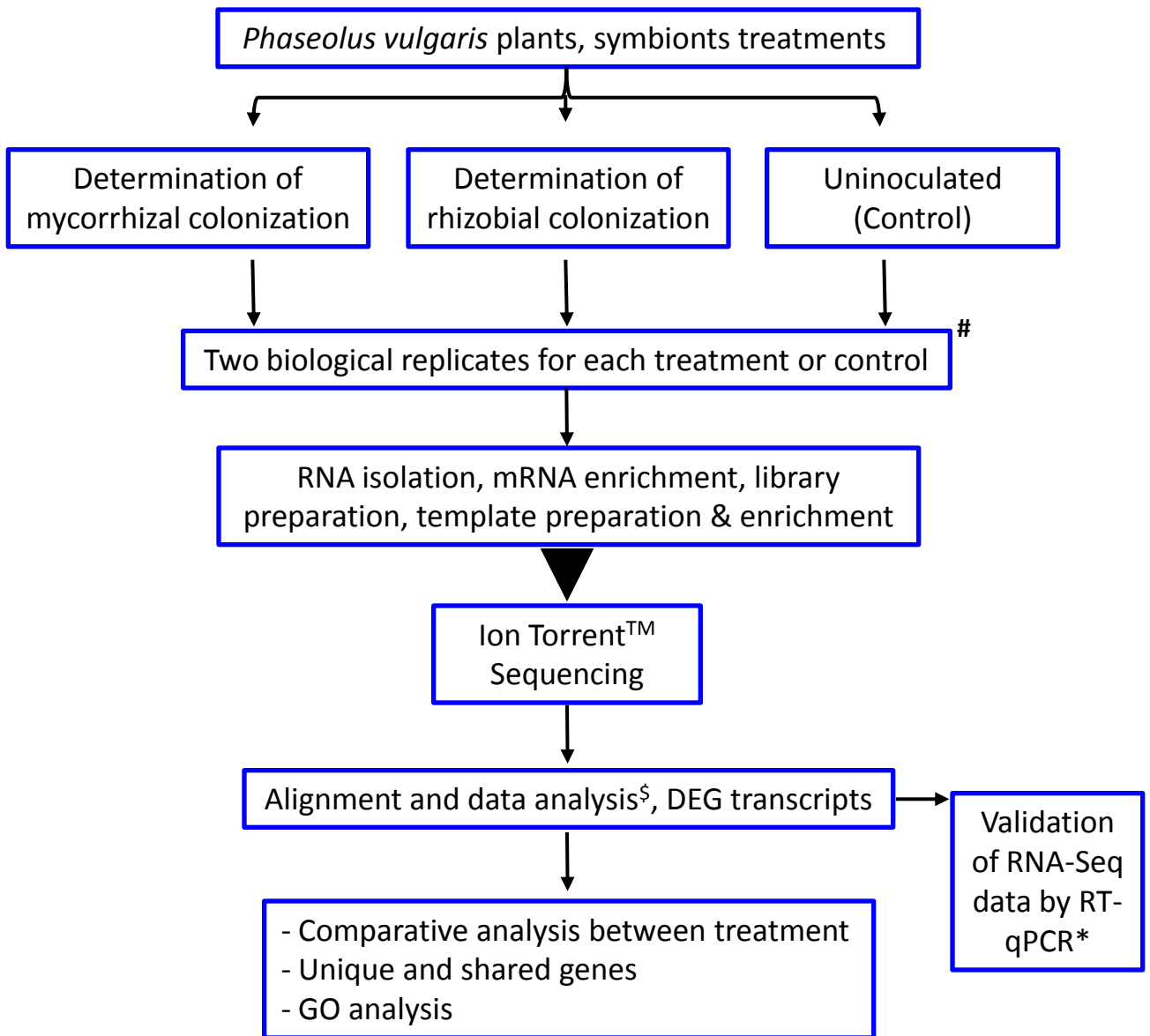


S1 Fig



S1 Fig. Overview of experimental design for RNA-sequencing and data analysis. Five-day-old wild-type *P. vulgaris* plants were inoculated with *Rhizophagus irregularis* for mycorrhizal colonization or *Rhizobium tropici* for the establishment of root nodules. At 2 weeks post inoculation (wpi), the symbiont-colonized roots were subjected to global RNA sequencing analysis. # Each biological replicate was a pool of 3 symbiont-colonized or control roots. \$ Data analysis includes normalization and quantification (fold-change in expression over control), and values were averaged for two replicates; * validation of RNA-Seq data by RT-qPCR from three biological replicates.