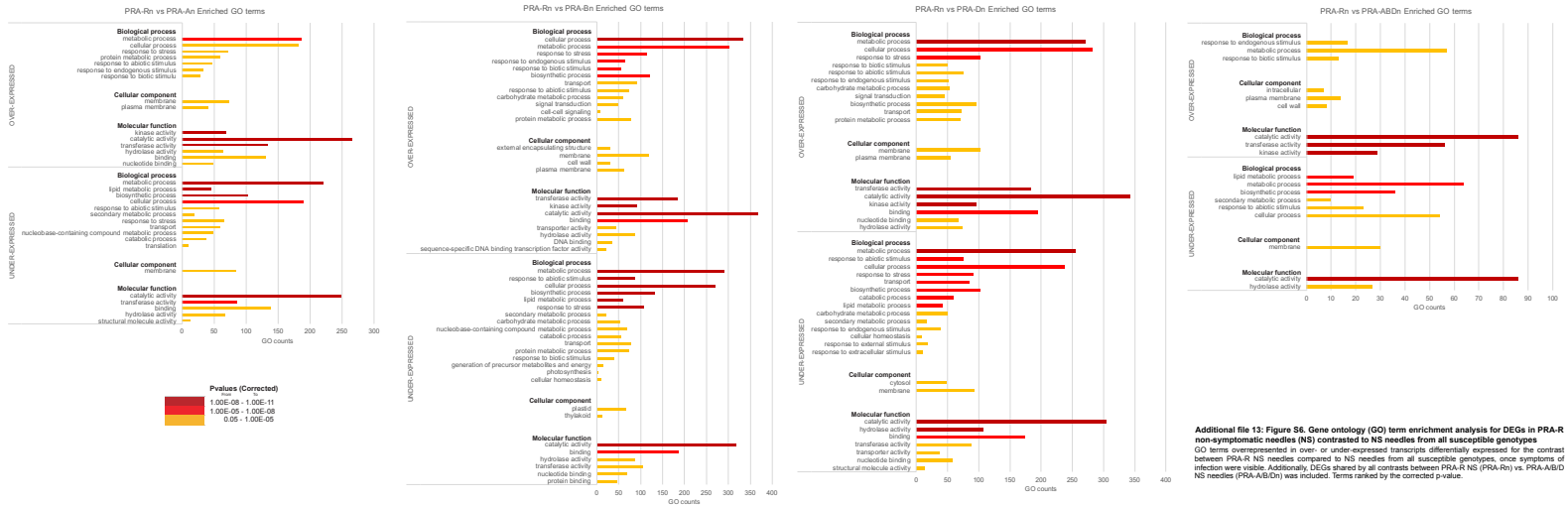




Additional file 13: Figure S6. Gene ontology (GO) term enrichment analysis for DEGs in PRA-R non-symptomatic needles (NS) contrasted to NS needles from all susceptible genotypes
GO terms overrepresented in over- or under-expressed transcripts differentially expressed for the contrast between PRA-R NS needles compared to NS needles from all susceptible genotypes, once symptoms of infection were visible. Additionally, DEGs shared by all contrasts between PRA-R NS (PRA-Rn) vs. PRA-AB/D NS needles (PRA-AB/Dn) was included. Terms ranked by the corrected p-value.