Figure S1. Root phenotype under control (first and third row) and acid soil (second and fourth row) for 1 day treatment (A), 3 days treatment (B).
**Figure S2.** Function analysis of genes differentially expressed in S1587-17 after 1 day of treatment in acid soil. A: Up-regulated; B: Down-regulated. All the genes that didn’t present Gene Ontologies were removed from the analysis.
Figure S3. Function analysis of genes differentially expressed in S1587-17 after 3 days of treatment in acid soil. A: Up-regulated; B: Down-regulated. All the genes that didn’t present Gene Ontologies were removed from the analysis.