Supplementary Figure 6: Blast2Go-derived combined graphs for the 26 annotated genomic regions hypothesized to be recurrently involved in divergent edaphic adaptation. Only categories with a node score ≥ 2.00 are shown. a) Most important GO biological processes, and b) the GO categories of molecular functions of those genomic regions. Nodes are coloured by node score (i.e., relevance).