



Figure 7. Visualization of proteomic variations in symptomatic areas (G7C) with respect to healthy leaves (G0). Distribution of DAPs in the most represented functional categories by MapMan analysis. Up- and down-regulated proteins are shown separately. Aminoacyl-tRNA synthetase, ARS; BCAA, branched-chain amino acid; Nt, nucleotide; OPPP, oxidative pentose phosphate pathway; PGL, pathogen polygalacturonase inhibitor; TCAC, tricarboxylic acid cycle; USP, universal stress protein.