



Figure 3. Metabolite profiling in curled and healthy leaves of *P. persica*. **A.** Heat map representing metabolite profiling performed by GC-MS. The heat map displays the relative level of each normalized metabolite to its amount found in G0 leaves (healthy leaves) and expressed as log₂. The color scale is shown at the top of the figure and it is proportional to the content of each metabolite. Grey color indicates that the metabolite was not detected, therefore there is not assigned value. At least five biological independent determinations were analyzed for each sample. Relative values for each metabolite peak area \pm SD are displayed in Table S1 and plotted with their corresponding static analysis in Figure S1. **B.** Principal component analysis (PCA) of metabolite composition by GC-MS. Between parentheses, the variance explained by each component (%) is shown. **C.** Sorbitol and starch contents enzymatically determined. Starch amount is expressed as nmols of glucose released per milligram of fresh weight. Bars with at least one same letter are not significant statically different.