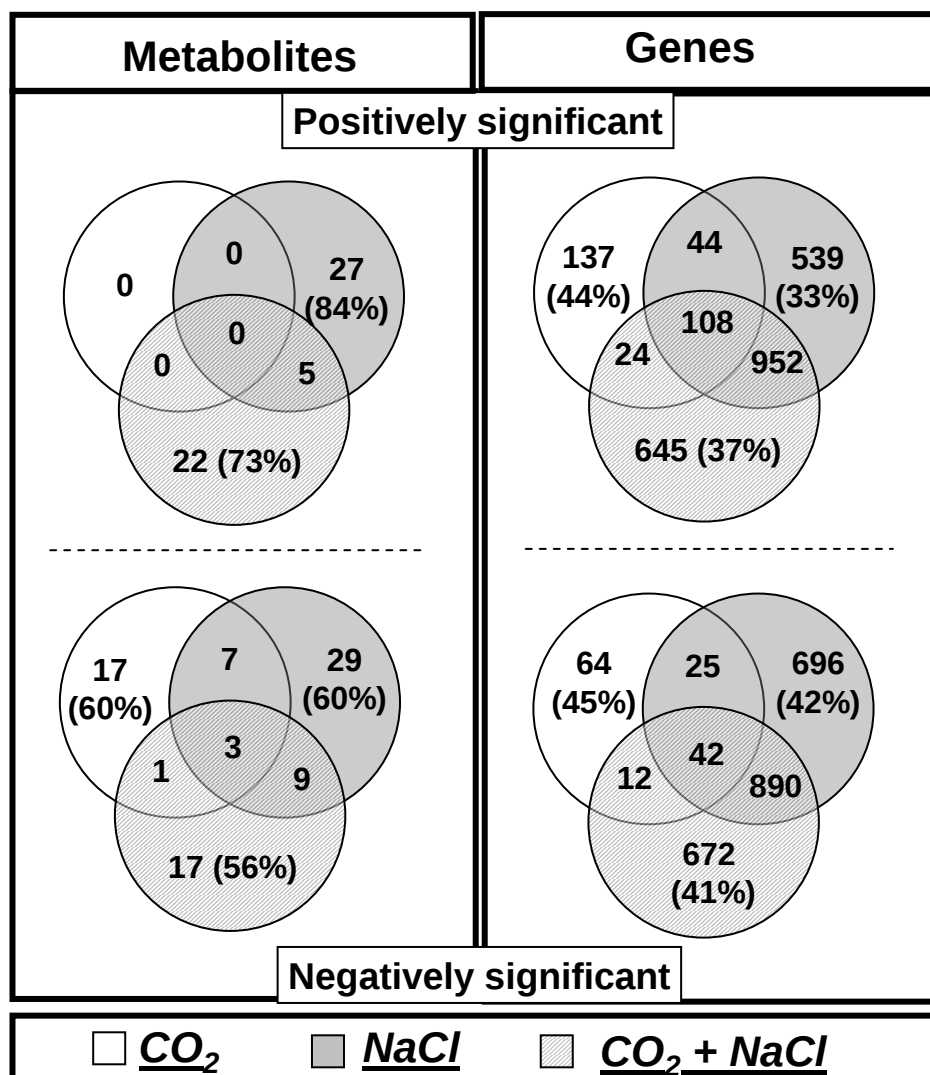


Additional File 3

Number of common positively and negatively significant genes and metabolites between the elevated CO₂, the NaCl and the combined stresses, as identified from paired-SAM.



The percentage in parenthesis indicates the fraction of the genes or metabolites that were identified as positively or negatively significant only under the particular stress in the total number of genes or metabolites that were used in the analysis of the particular stress.