

Table S1: Cell wall over-expressed genes and proteins in AC-DC compared to roots in control condition, related to figure 2.
 See excel file, sheet “Table S1”. The column “Bin name” corresponds to gene ontologies obtained with Mapman. ID = identifier. Bold identifiers correspond to proteins. (fold change $\geq 1,5$; p-value $\leq 0,05$)

Table S2: Differentially expressed genes and proteins upon PEP13 elicitation in roots, related to figure 3.
 See excel file, sheet “Roots E vs NE”. (fold change $\geq 1,5$; p-value $\leq 0,05$)

Table S3: Differentially expressed genes and proteins upon PEP13 elicitation in AC-DC, related to figure 3.
 See excel file, sheet “AC-DC E vs NE”. (fold change $\geq 1,5$; p-value $\leq 0,05$)

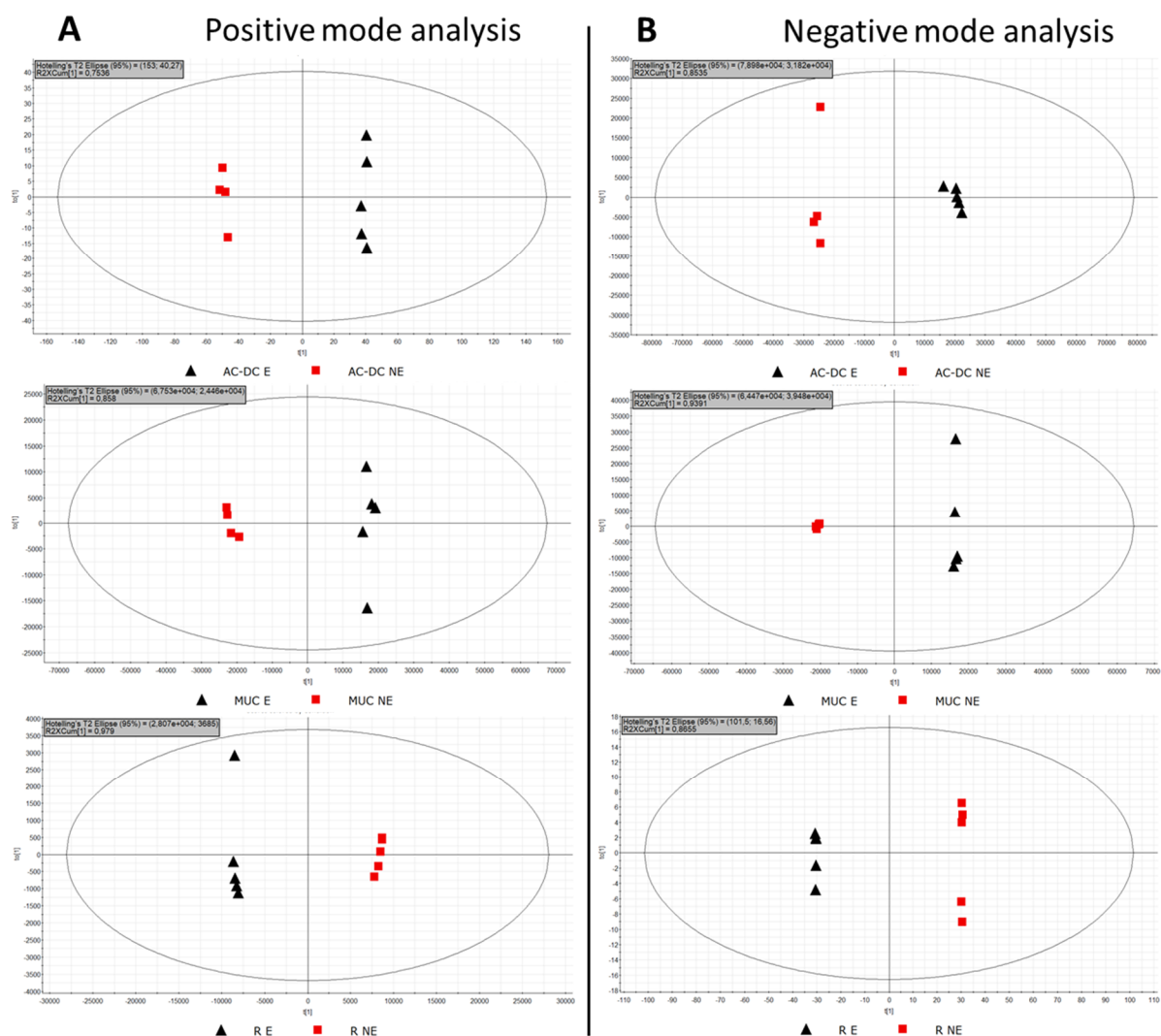


Figure S1: Supervised multivariate classification OPLS-DA of AC-DC, mucilage and roots with and without PEP-13 elicitation, related to metabolomic analysis.
 R =roots; MUC = mucilage; AC-DC = root Associated Cap-Derived Cells.