



Supplementary Figure S1. Unsupervised hierarchical cluster analysis carried out from UHPLC-ESI/QTOF-MS metabolomic analysis of leaves (A) and root exudates (B) of tomato plants under low salt or high salt conditions, treated with either protein hydrolysates-based biostimulants (PHs), or with PGPR *A. brasilense*. The fold-change based heat map was used to build hierarchical clusters (linkage rule: Ward; distance: Euclidean).