



S2 Fig. Additional metabolites detected by the LC-MS analysis. The size of the bars represents the total abundance, and coloured parts indicate ¹³C labelling as depicted in the legend. The samples are from WT cells, WT grown in ¹³C-glycerol, THT1/THT2 RNAi grown in ¹²C-glycerol and THT1/THT2 RNAi grown in ¹³C-glycerol. Natural abundance of ¹³C is 1 %, hence the labelling in ‘un-labelled’ samples.
* the identity of these metabolites was confirmed using a match with a standard.