



Figure S2. Volcano plots of TCR stimulated cells compared to unstimulated cells.

The \log_2 foldchange (60 min/untreated) for individual proteins is plotted against the DeqMS calculated p-value for each individual compartment (Cytosol, Nucleus and Membrane). Proteins that fulfilled the criteria for translocation (opposite direction of change in at least 2 compartments with an absolute \log_2 foldchange >0.201 as well as a p-value <0.05 in both) are highlighted and color coded based on the translocation as denoted above.