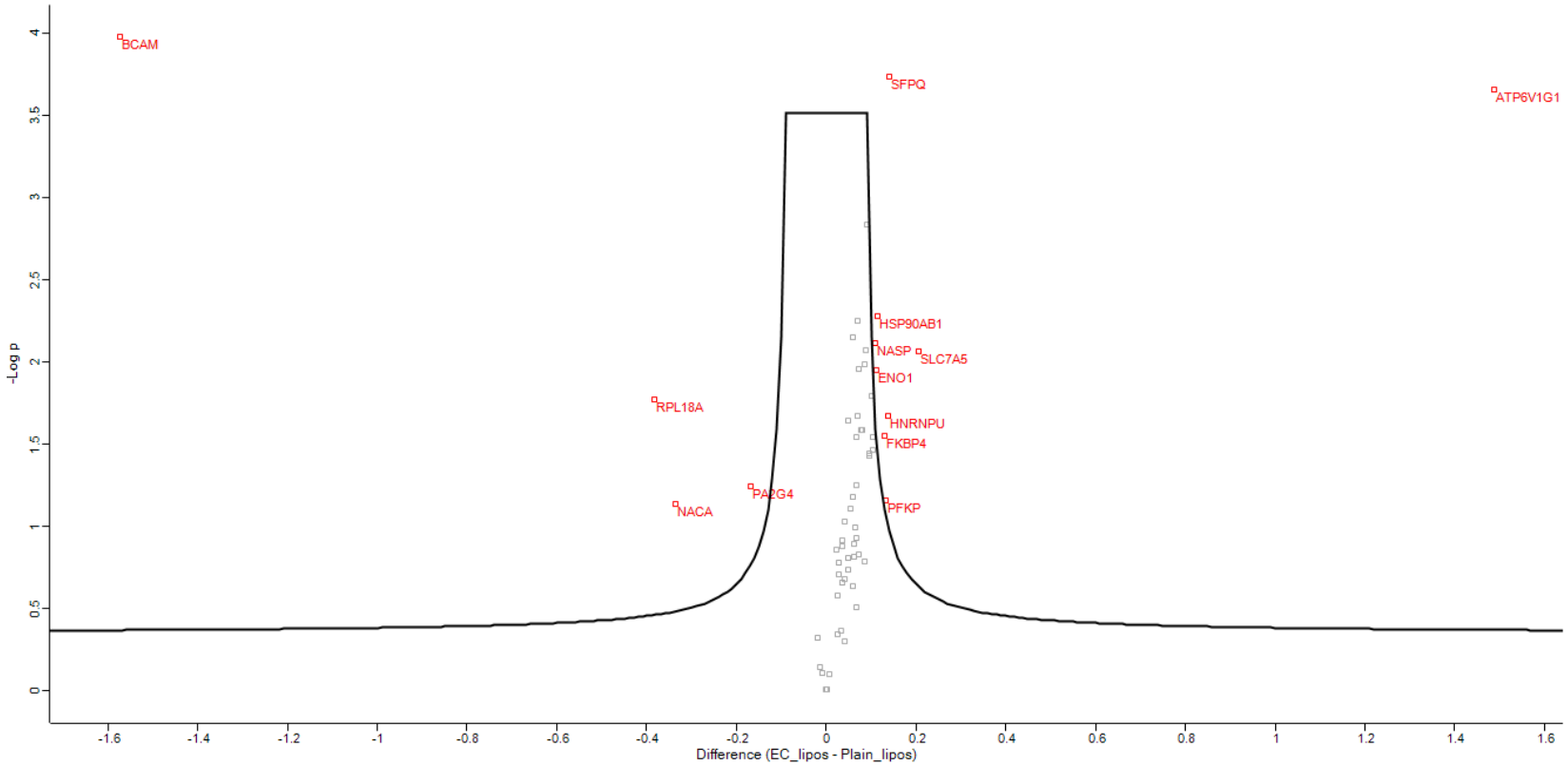


Figure S3. Volcano plot analysis of EC-labelled vs. plain SILAC liposomes conditions. Volcano plot displays fold-change differences in log2 heavy-to-light amino acid ratios between EC-labelled and plain SILAC liposomes conditions. Analysis includes all proteins identified as statistically significant in a one-way ANOVA test. X-axis represents the log2 fold-change in heavy-to-light amino acid ratio, Y-axis represents statistical significance as -log *p*-values. Analysis parameters: $S_0 = 0.1$, false discovery rate = 0.05. Significantly differentially incorporated proteins (red) are labelled with gene symbols. The volcano plot was generated using Perseus software.