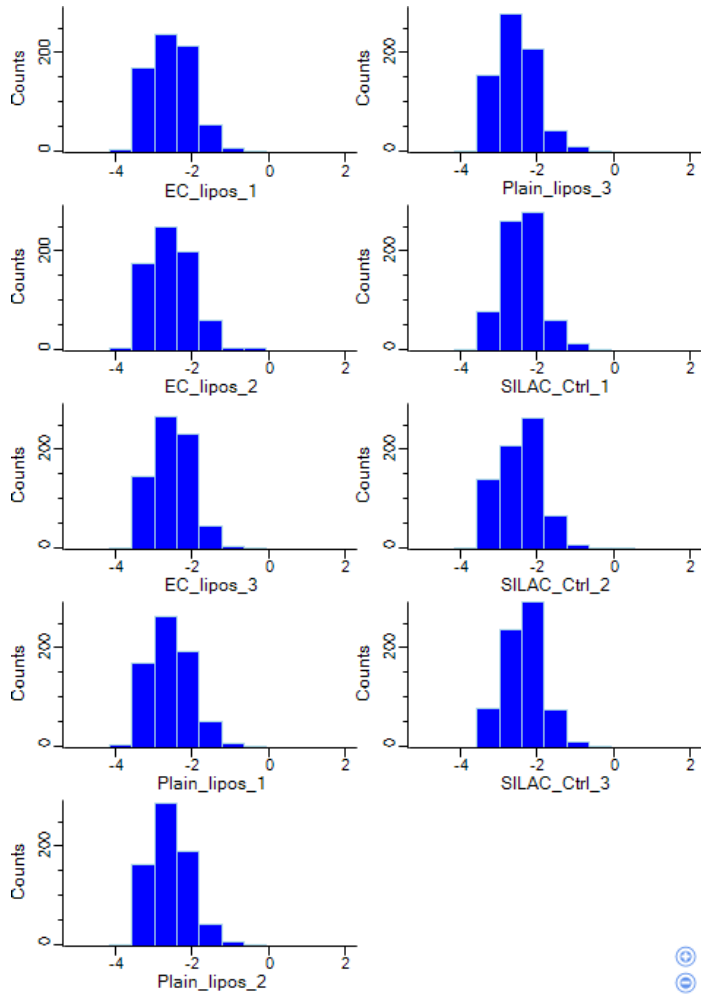


Figure S1. Histograms of EC-labelled SILAC liposomes, plain SILAC liposomes, and SILAC medium control triplicates. Heavy-to-light amino acid ratio data for 711 proteins across triplicate samples in each experimental condition were visualised as histograms to evaluate whether the data followed a normal distribution after log2 transformation. X-axis represents the log2 heavy-to-light ratios, Y-axis indicates their frequency. Histograms were generated using Perseus software.