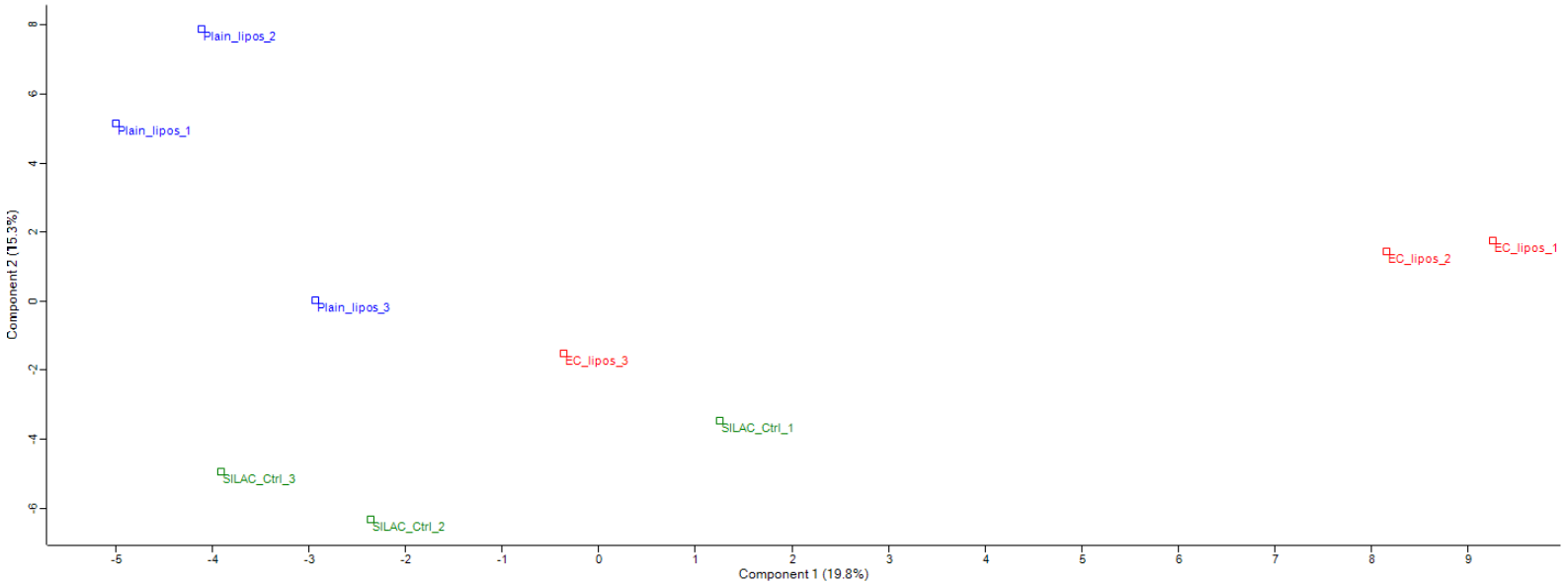


Figure S2. Principal component analysis (PCA) of EC-labelled SILAC liposomes, plain SILAC liposomes, and SILAC medium control triplicates. PCA was performed on heavy-to-light ratio data for 711 proteins across triplicate samples in each experimental condition to assess data similarity between replicates and differences between treatment groups. X-axis represents Principal Component 1, which accounts for the greatest variance in the dataset. Y-axis represents Principal Component 2, which accounts for the second greatest variance in the dataset. EC-labelled SILAC liposomes triplicates are highlighted in red, plain SILAC liposomes in blue, and SILAC medium controls in green. The PCA plot was generated using Perseus software.