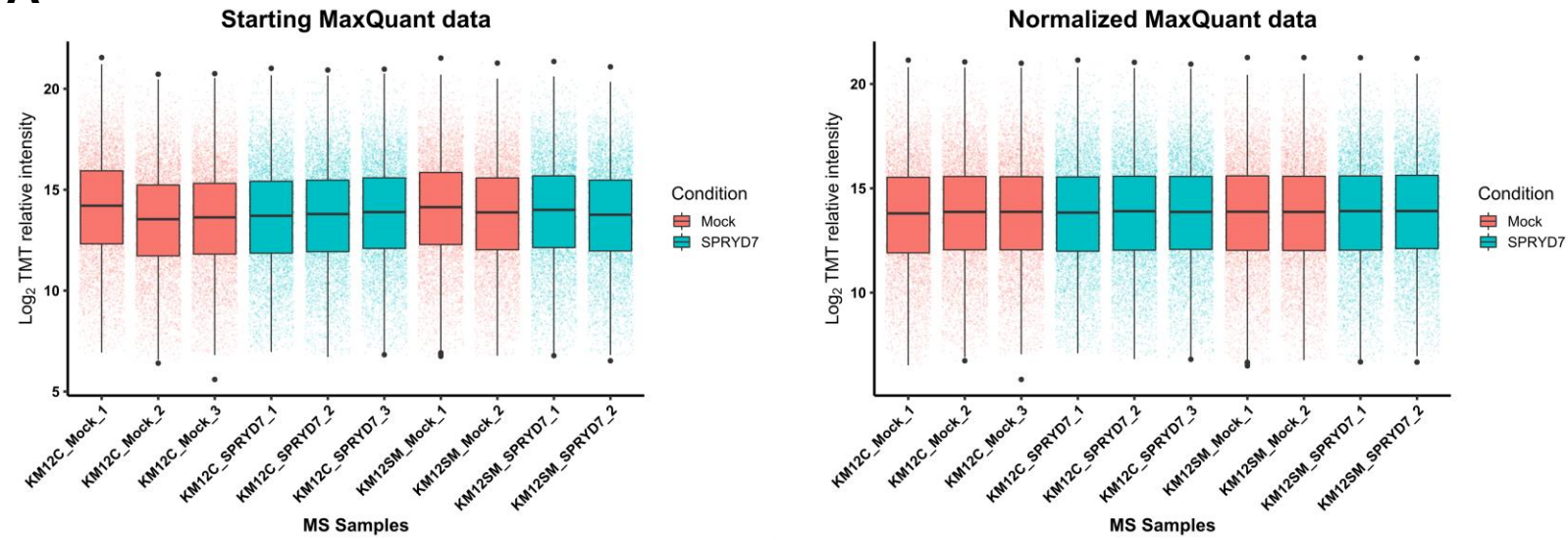
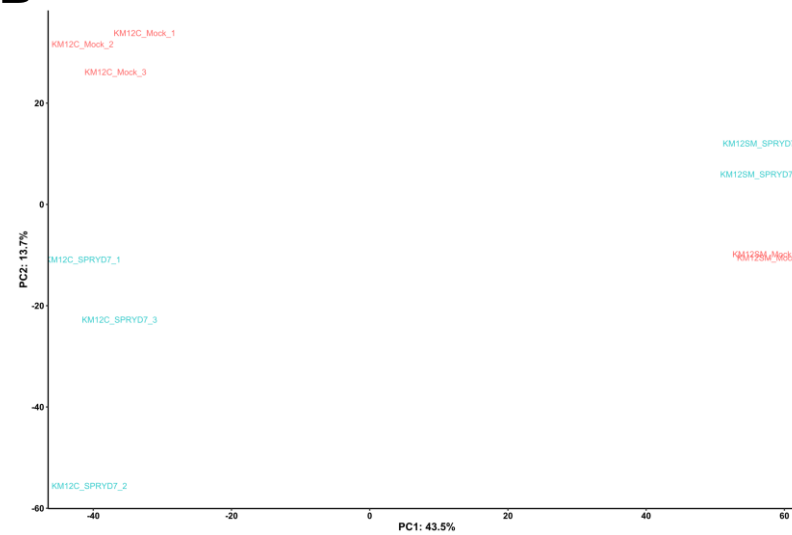


A**B**

Supplementary Figure 2. Quality control of the mass spectrometry data obtained from the TMT analysis. **(A)** SL normalization of MaxQuant data was performed to equal the total sum of protein intensities per TMT channel. **(B)** Principal component analysis (PCA) demonstrate the appropriate clustering of biological replicates according to their biology.

Supplementary Figure 2