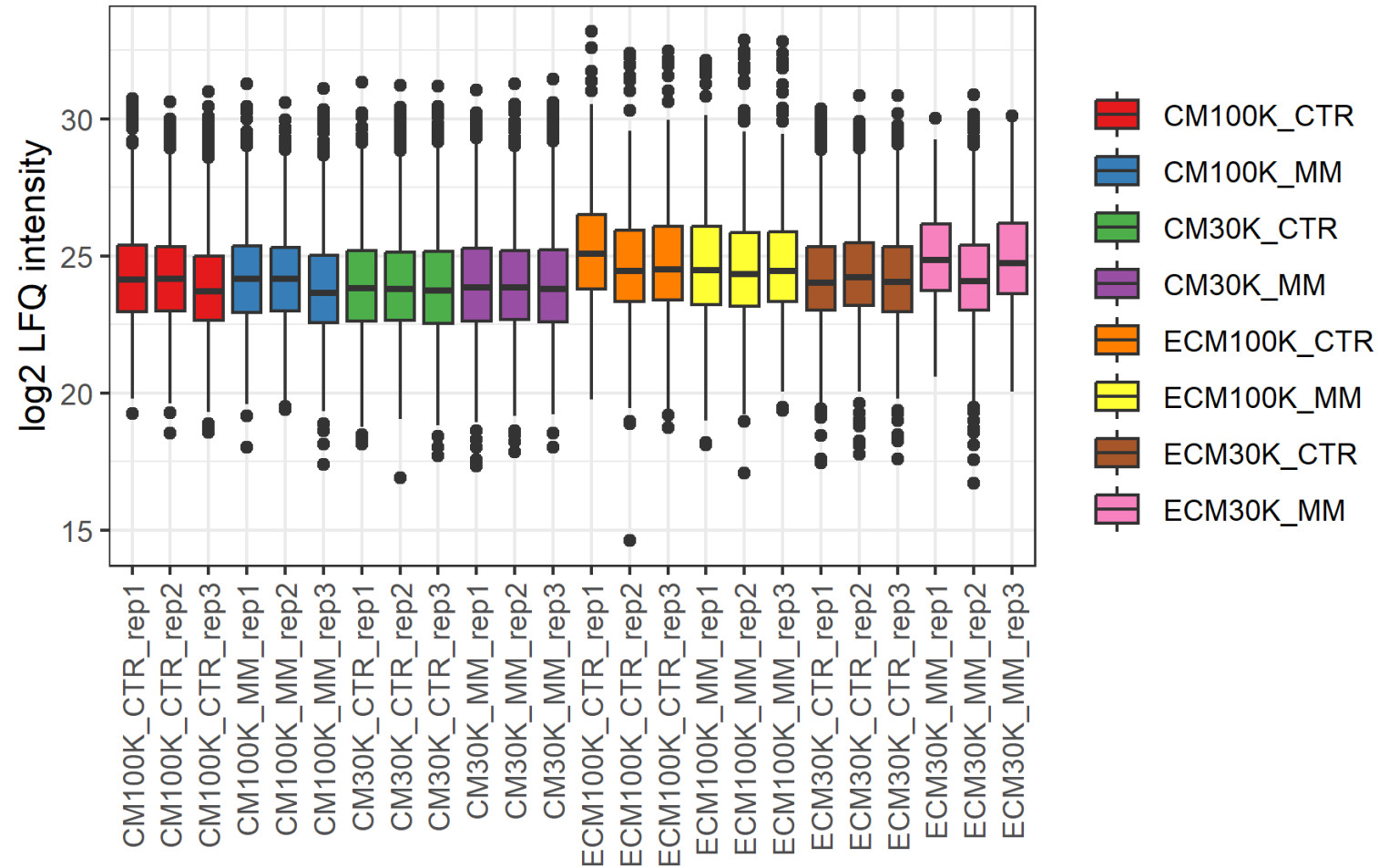
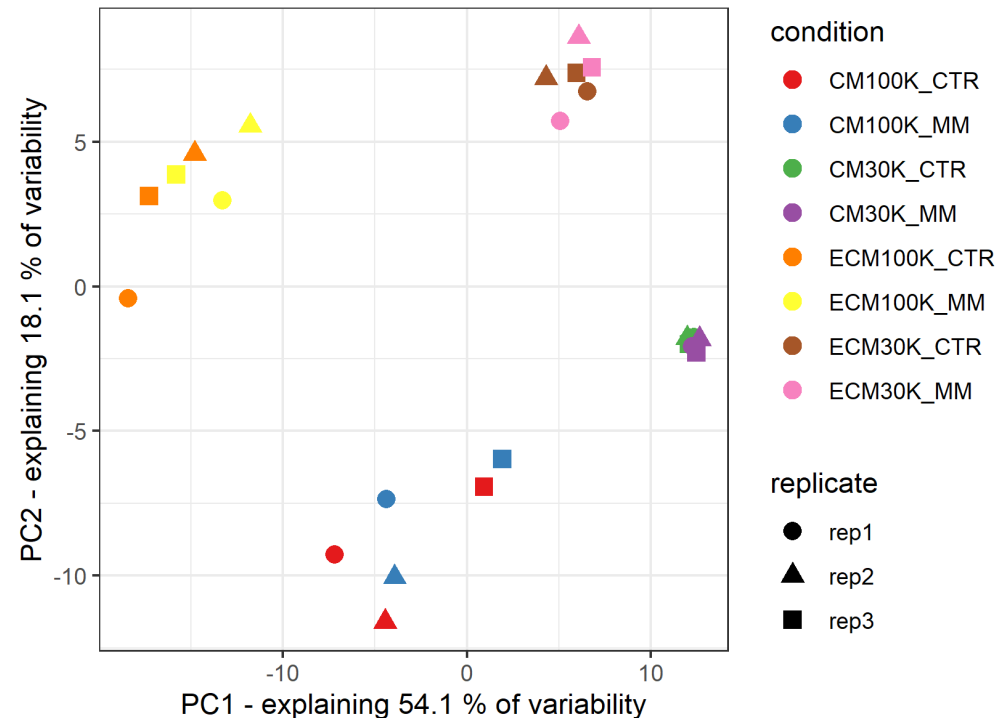


Samples main feature intensity distributions



Principal component analysis (PCA)

A PCA was performed on the replicate samples using all quantified proteins as variables. The scatter plot below shows sample mappings along the two principal components (PCs). Percentages of explained data variance for each PC are shown on the x and y axis.



Heatmap of between-samples correlations

A heatmap of sample correlations after hierarchical clustering is shown below. Plotted values are Spearman correlation coefficients between protein expression values of all pairs of samples.

