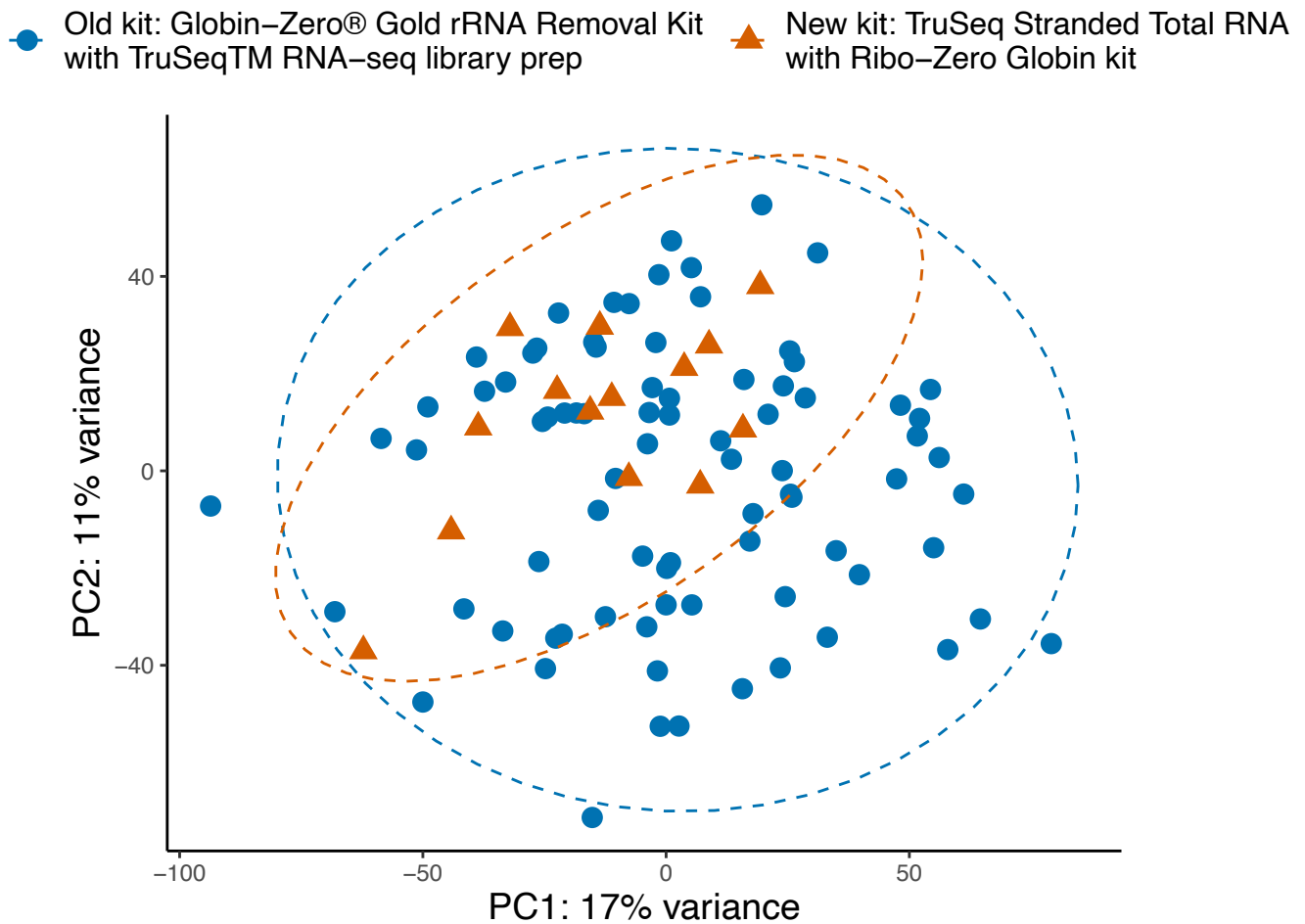




Supplementary Figure 3: Principal component analysis plot of the gene expression count data, demonstrating the variation between the samples. Each dot represents a sample, the colors/shapes represent the version of depletion kit used during library preparation. The counts are transformed using the `varianceStabilizingTransformation` function in DESeq2. PC = Principal component.