Figure 1: Barplot showing classification accuracy on batch corrected TCGA RNA-seq data. Note that for un-batch corrected RNA-seq data, classification accuracy is slightly lower than for the batch corrected data (3,029 compared to 3,136; \( P = 1.5 \times 10^{-3} \)). This suggests that the batch correction procedures were effective. See Methods for description of batch correction procedure.