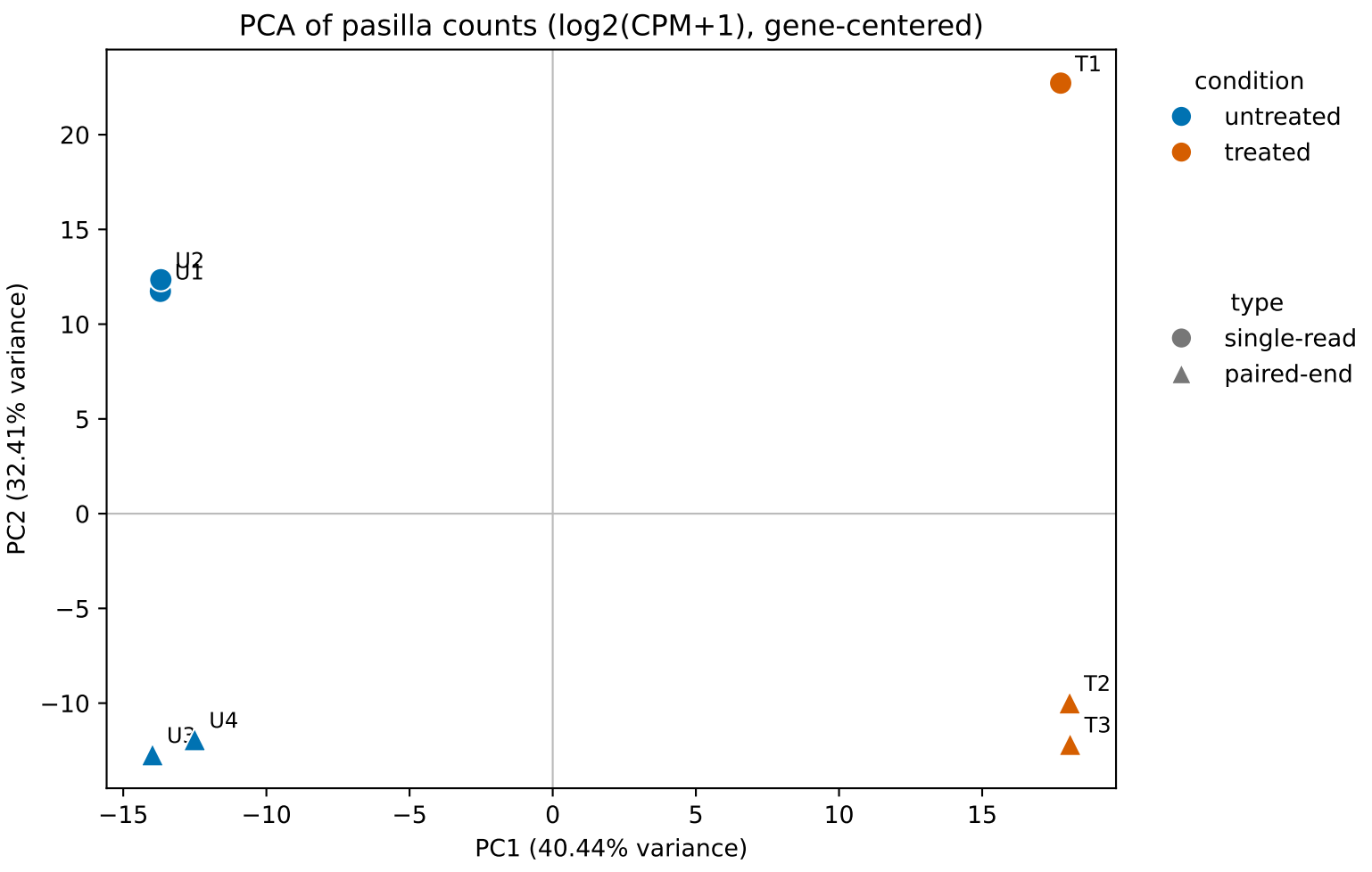

PCA of pasilla counts (log2(CPM+1), gene-centered)
PC1 (40.44% variance)
PC2 (32.41% variance)
condition
untreated
treated
type
single-read
paired-end
T1
T2
T3
U1
U2
U3
U4