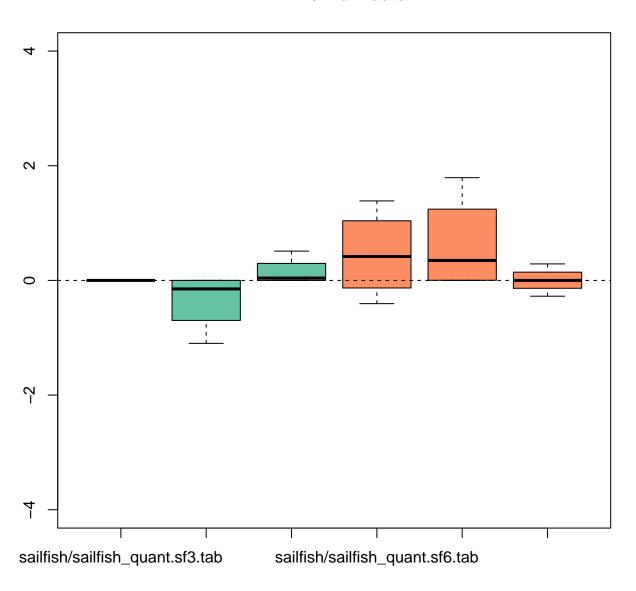
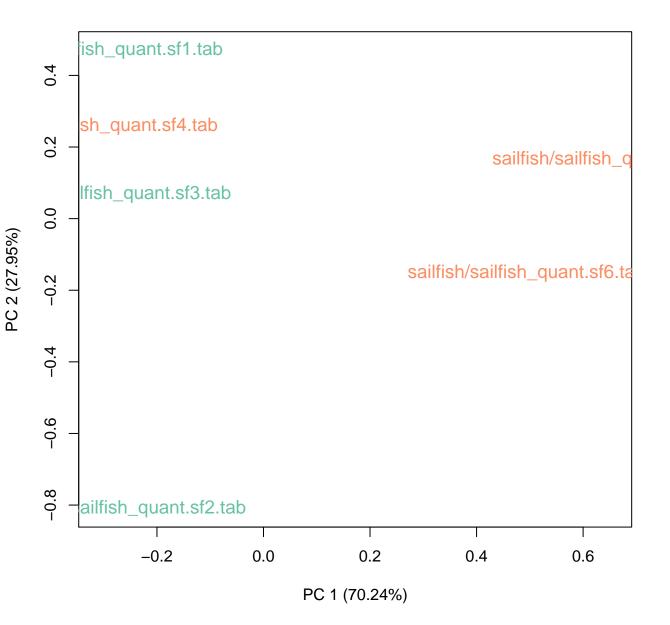
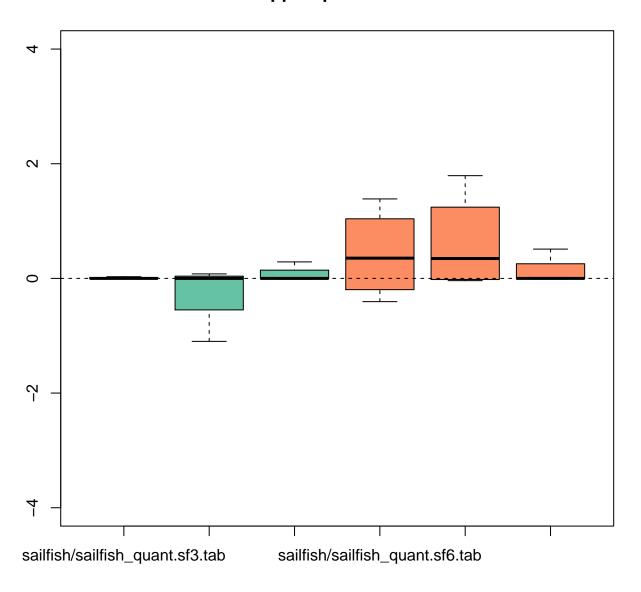
RLE for raw data



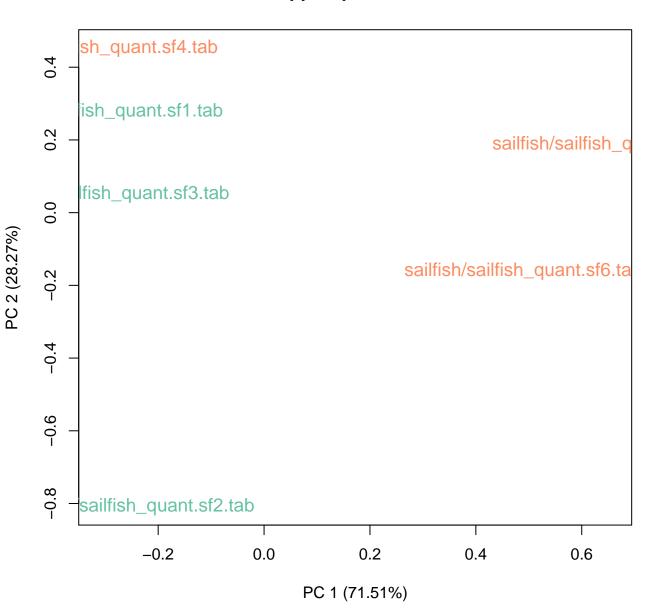
PCA for raw data



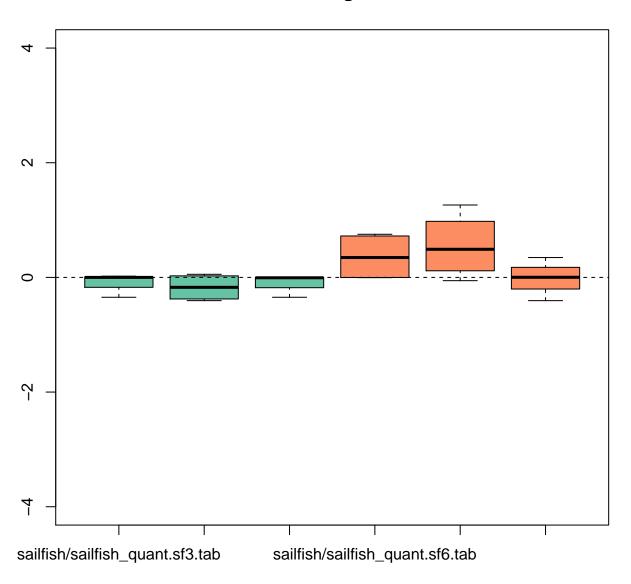
RLE for upper quartile normalized



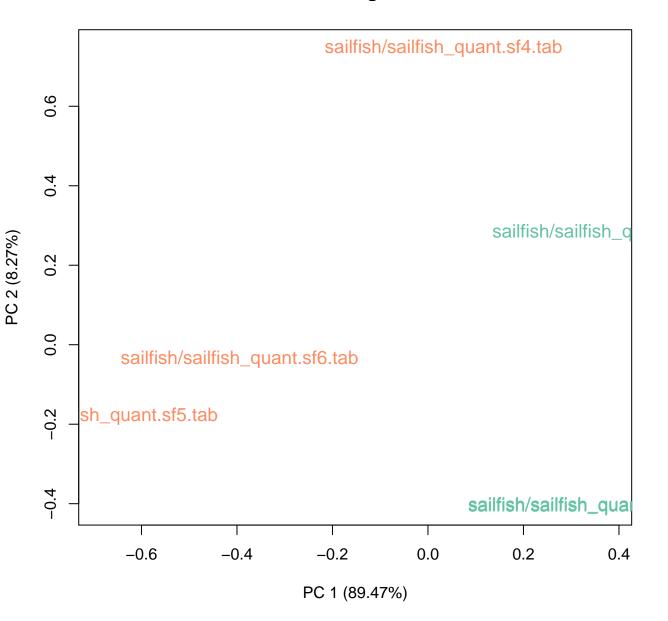
PCA for upper quartile normalized



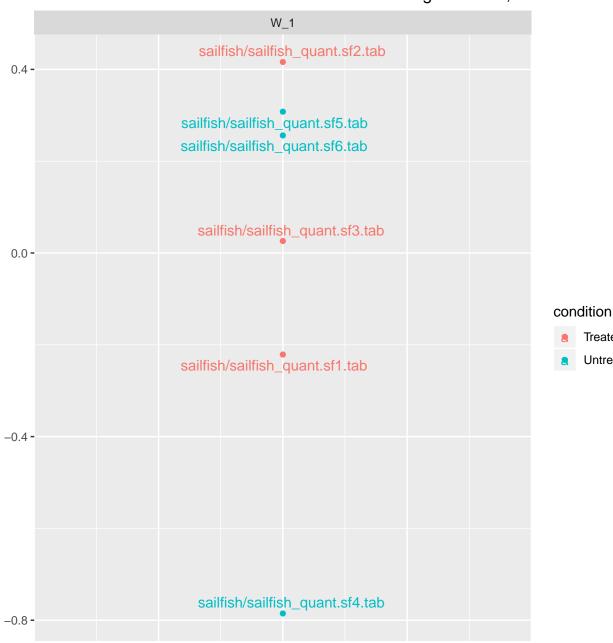
RLE for RUVr using residuals, k=1



PCA for RUVr using residuals, k=1

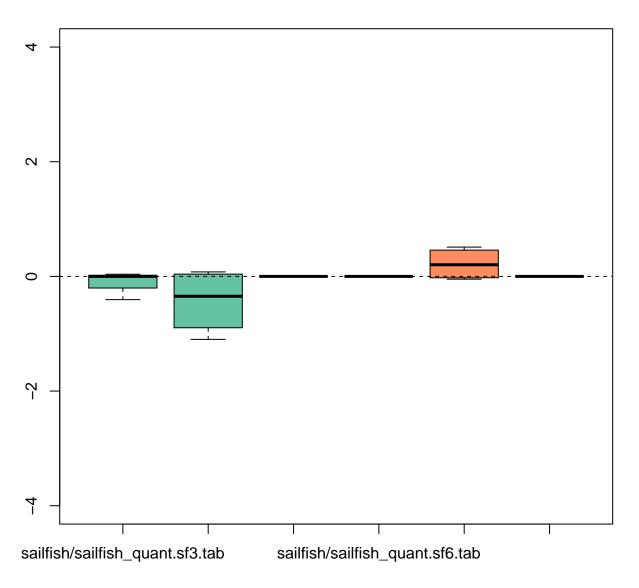


Factors of unwanted variation for method: RUVr using residuals, k=1

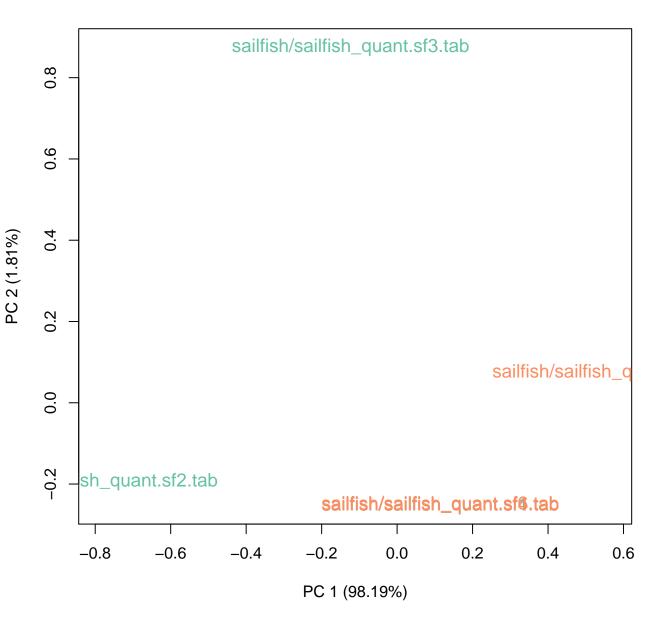


Treated Untreated

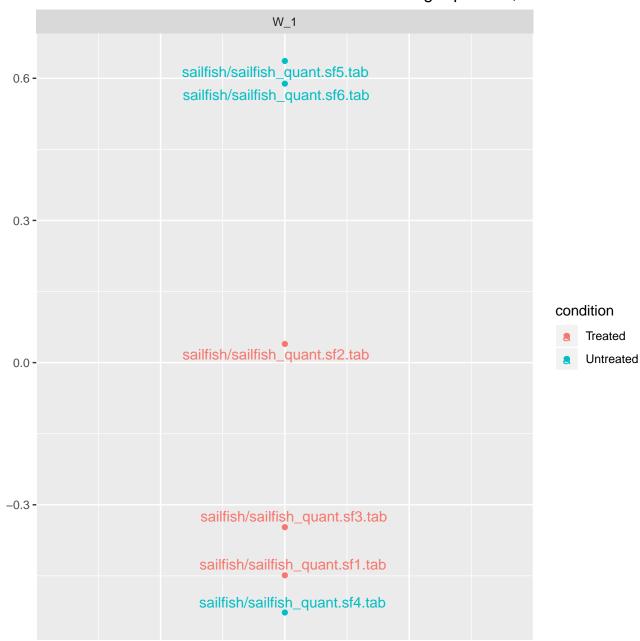
RLE for RUVs with replicate samples, k=1



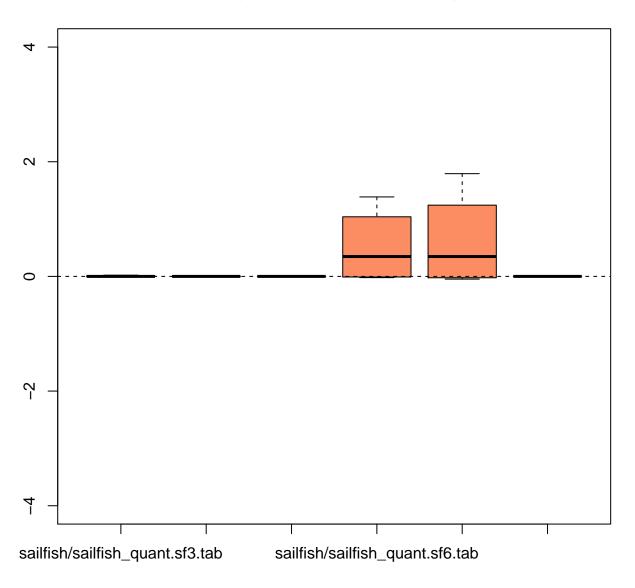
PCA for RUVs with replicate samples, k=1



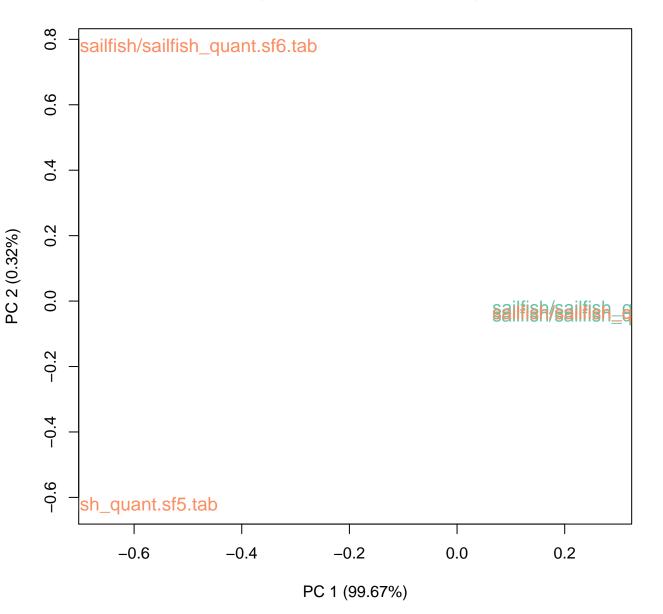
Factors of unwanted variation for method: RUVs using replicates, k=1



RLE for RUVg with empirical control genes, k=1



PCA for RUVg with empirical control genes, k=1



actors of unwanted variation for method: RUVg with empirical control genes, k=1 W_1 sailfish/sailfish_quant.sf2.tab 0.5 sailfish/sailfish_quant.sf6.tab condition Treated Untreated 0.0 sailfish/sailfish_quant.sf3.tab sailfish/sailfish_quant.sf5.tab sailfish/sailfish_quant.sf1.tab -0.5 sailfish/sailfish_quant.sf4.tab