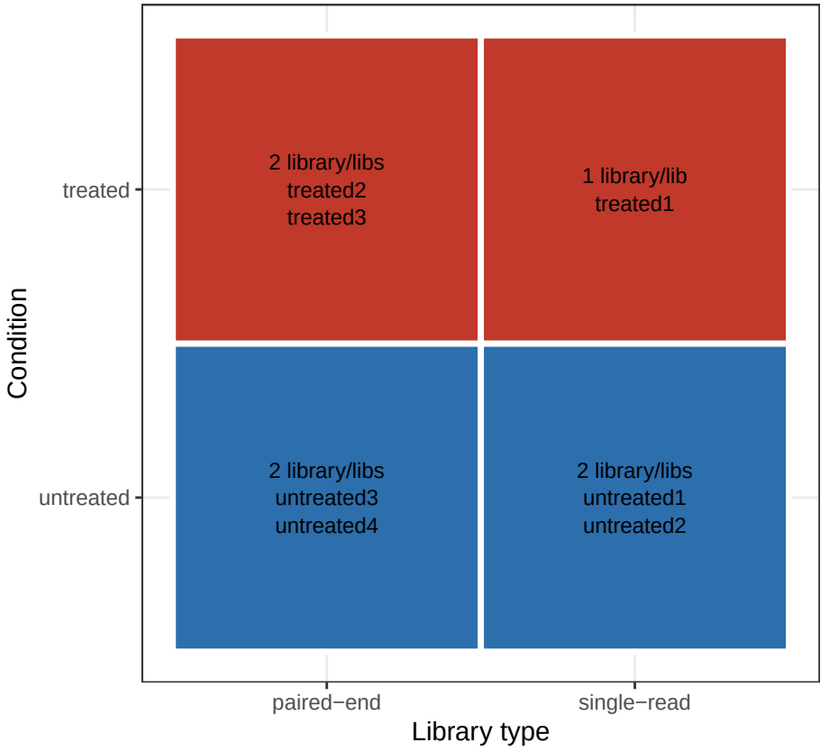
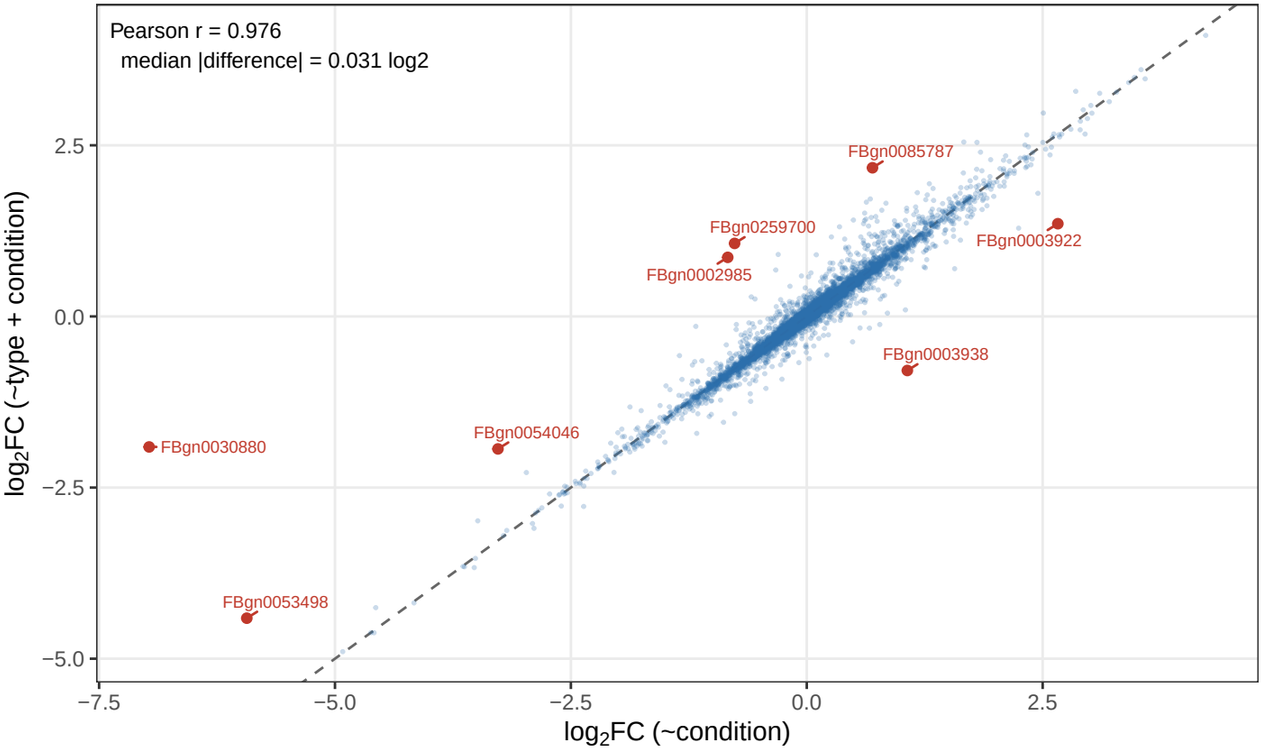
Design cells: condition x sequencing type

Numbers are independent experimental units (libraries)



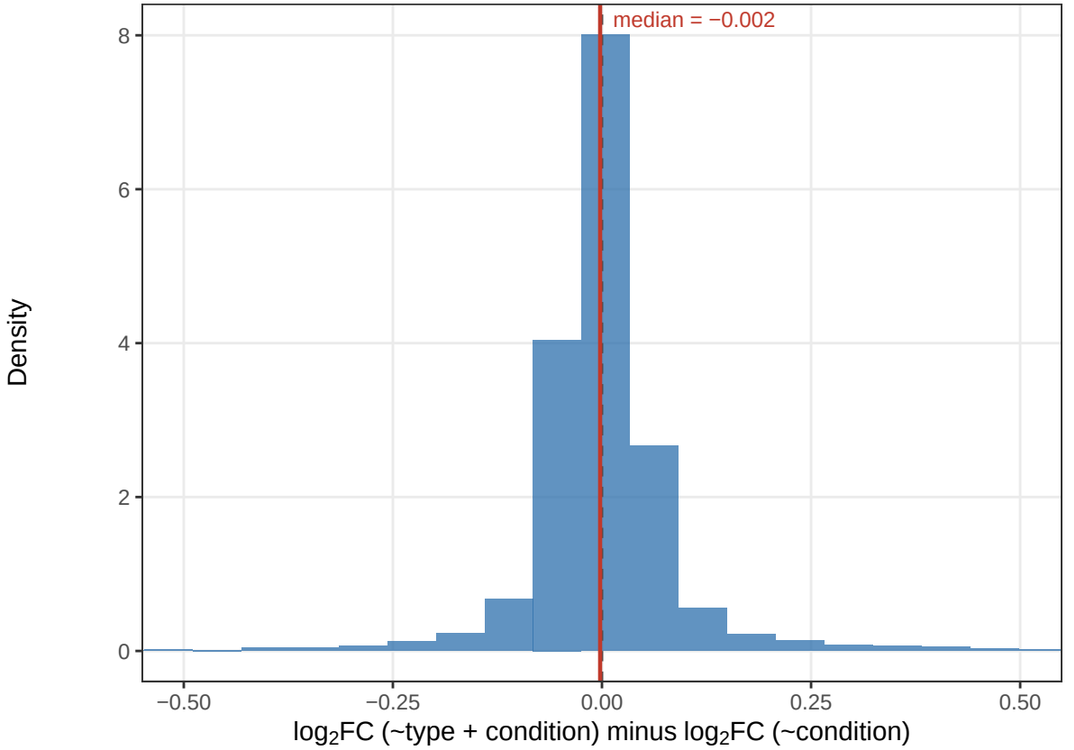
Treatment effect with vs without type adjustment

Unshrunken DESeq2 log2FC, treated vs untreated; dashed line y = x



Per−gene difference in treatment effect

x−axis truncated to [−0.5, 0.5]; 0.9% of genes outside



Effect difference vs mean expression

y−axis truncated to [−0.8, 0.8]; points beyond are not shown

