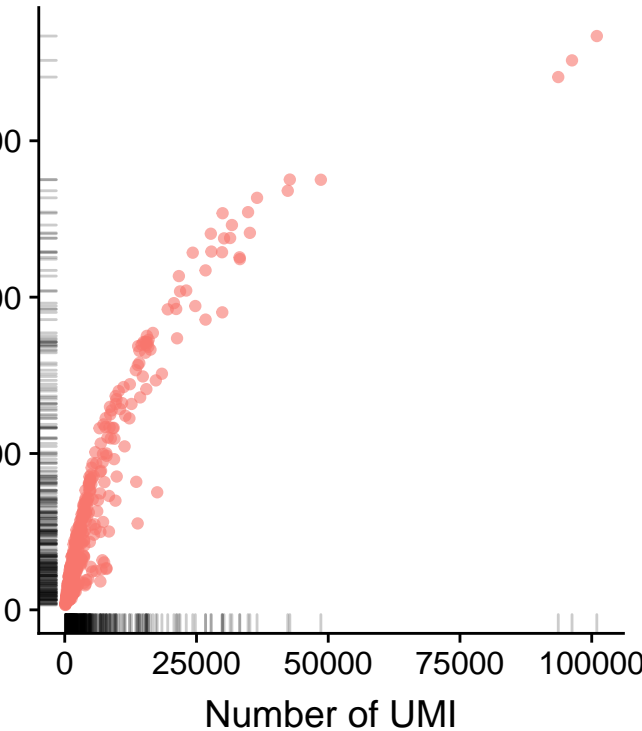
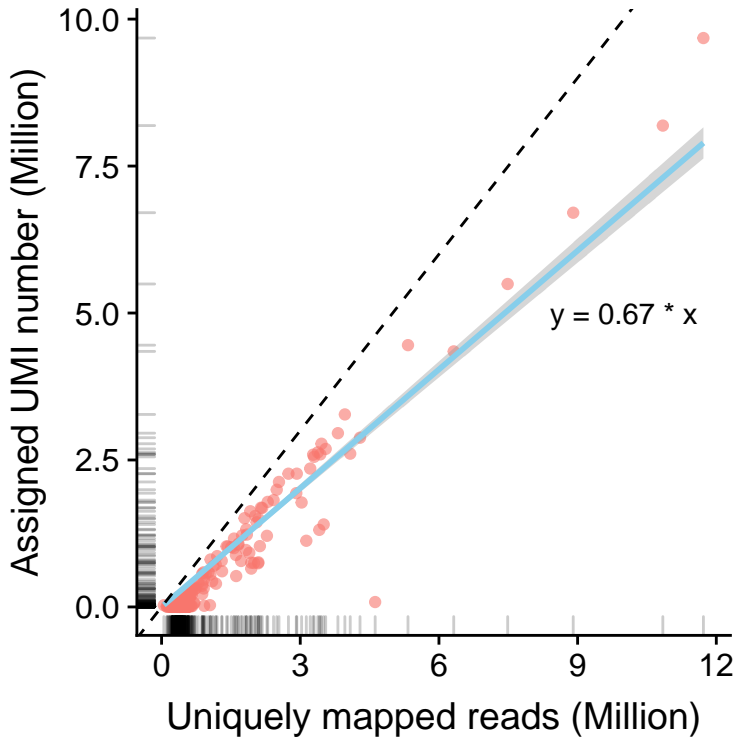
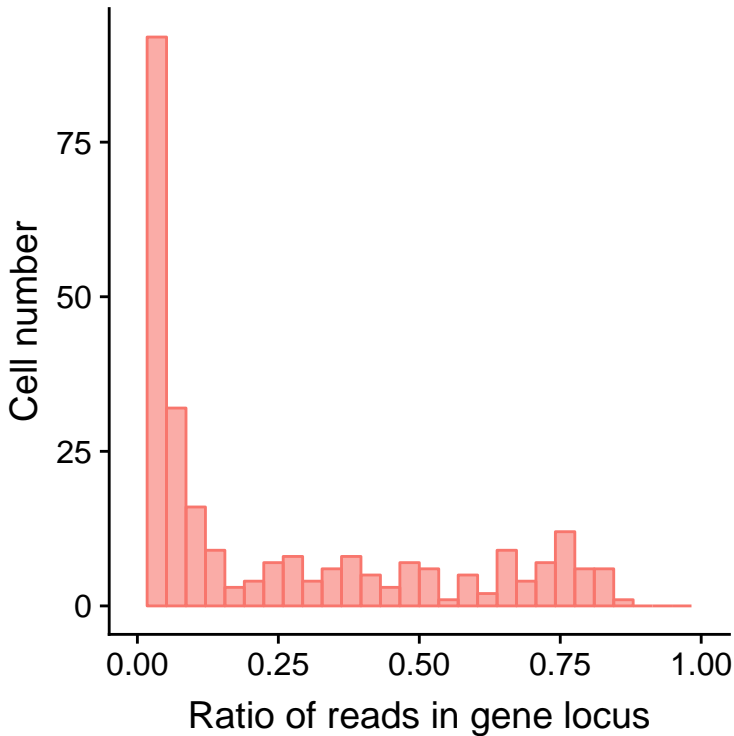


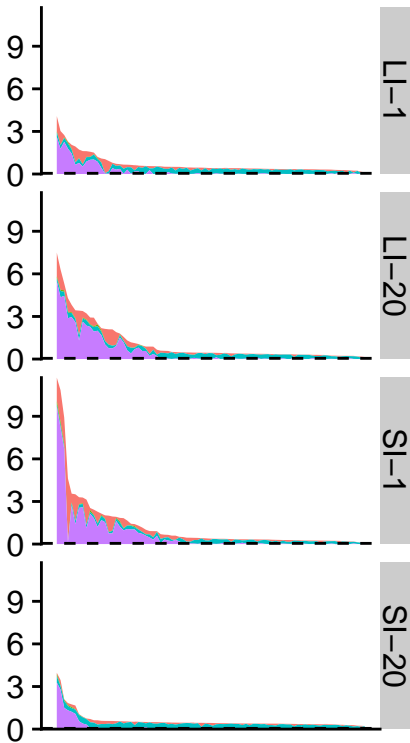
Number of detected genes







Uniquely mapped reads (Million)



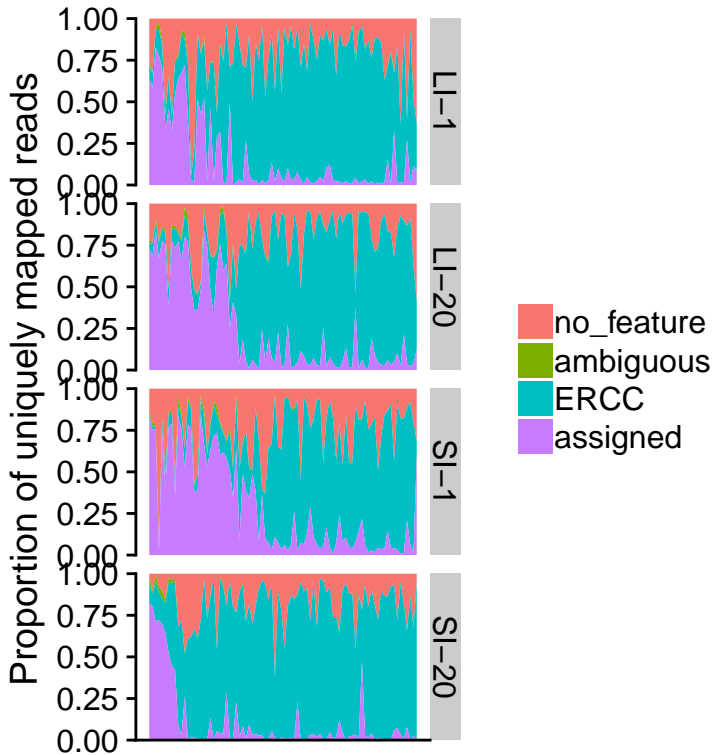
LI-1

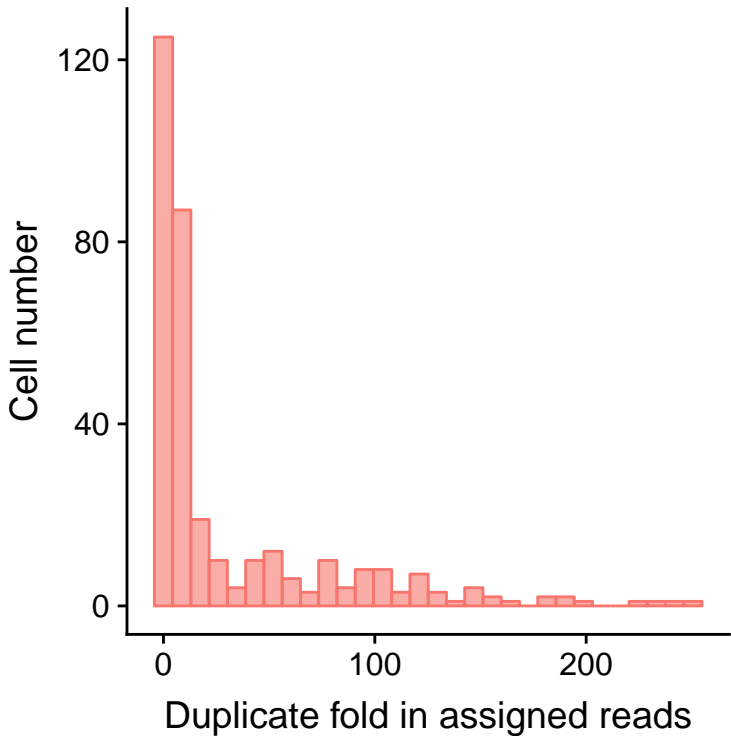
LI-20

SI-1

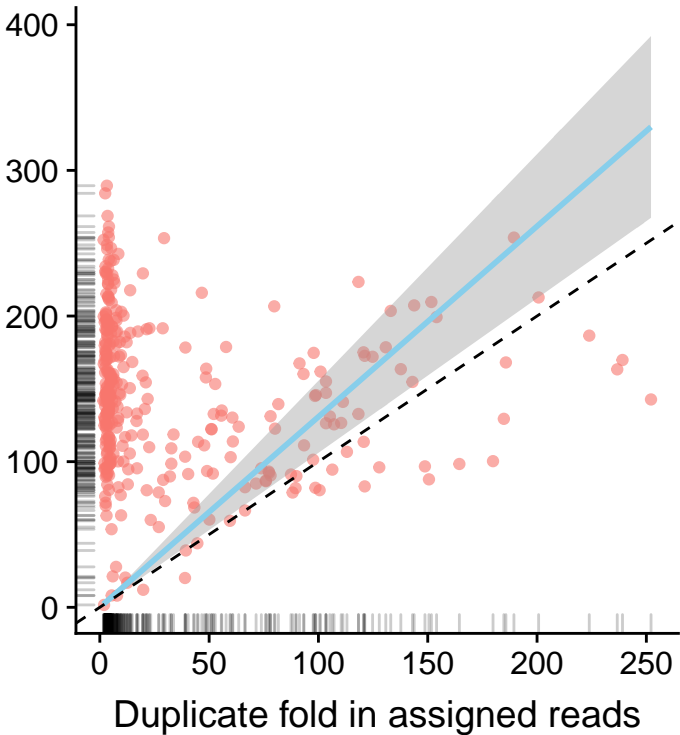
SI-20

no_feature
ambiguous
ERCC
assigned



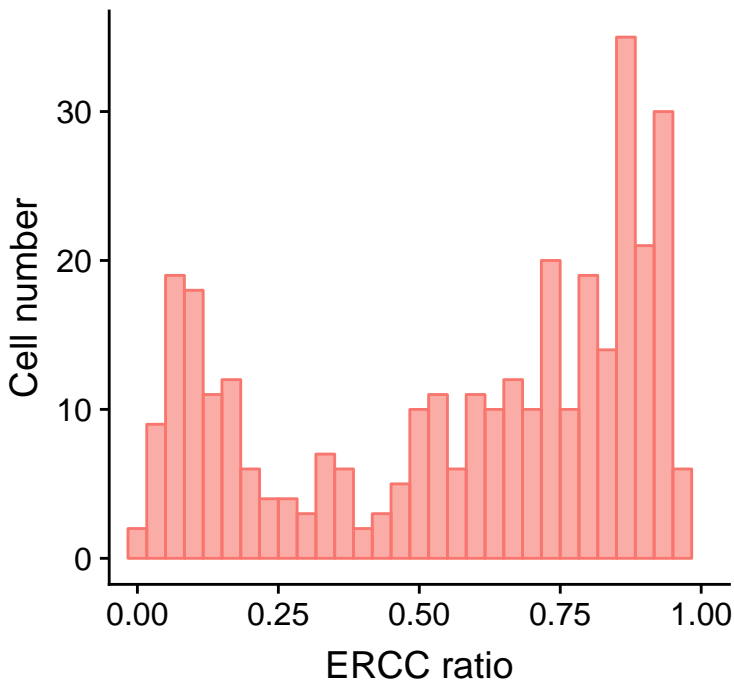


Duplicate fold in ERCC reads

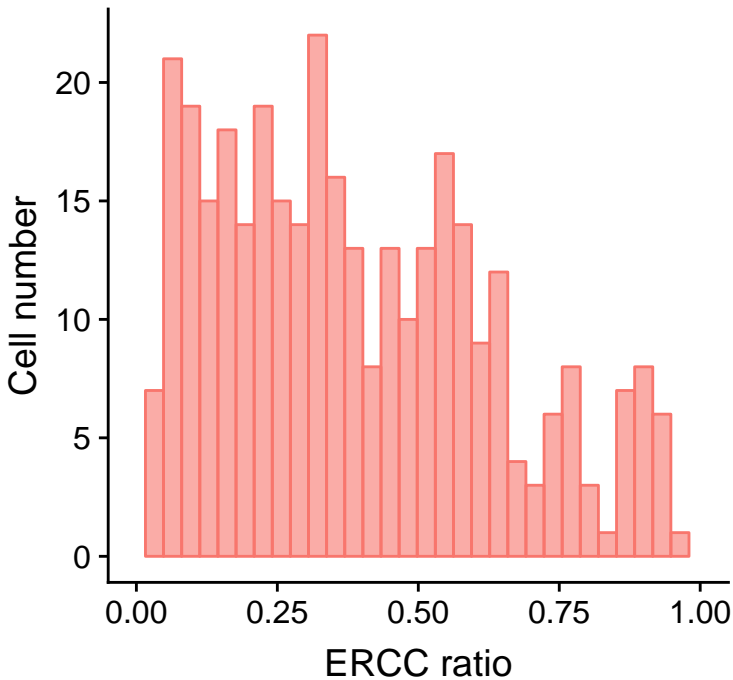


Duplicate fold in assigned reads

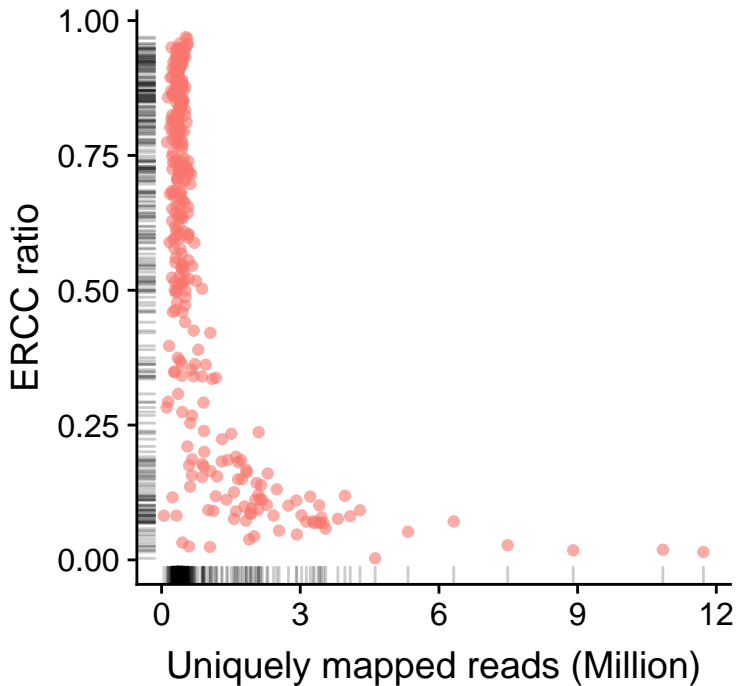
Before removing duplicates



After removing duplicates



Before removing duplicates



After removing duplicates

