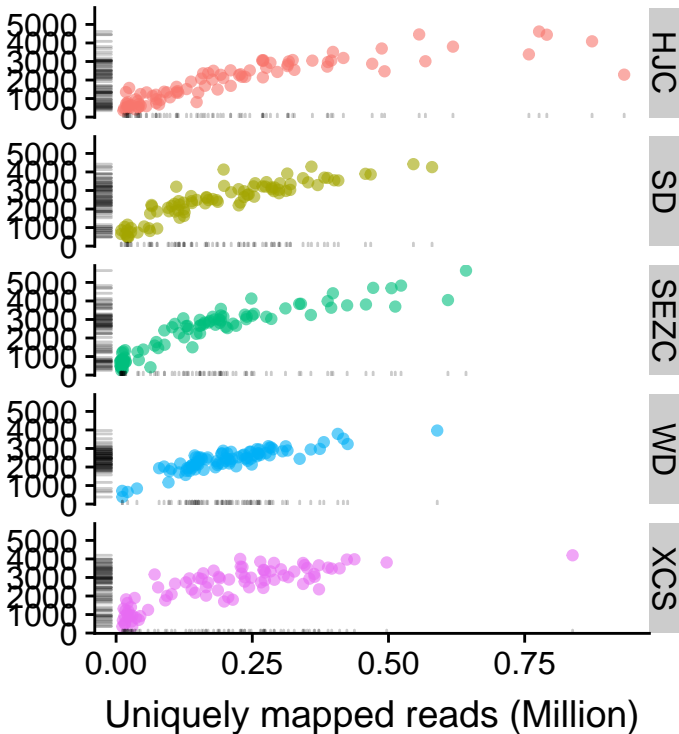
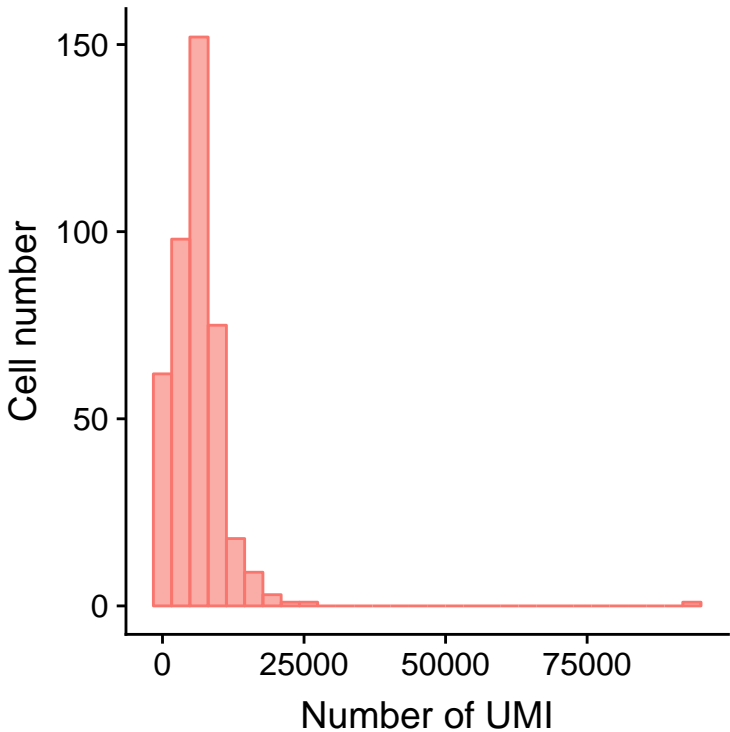
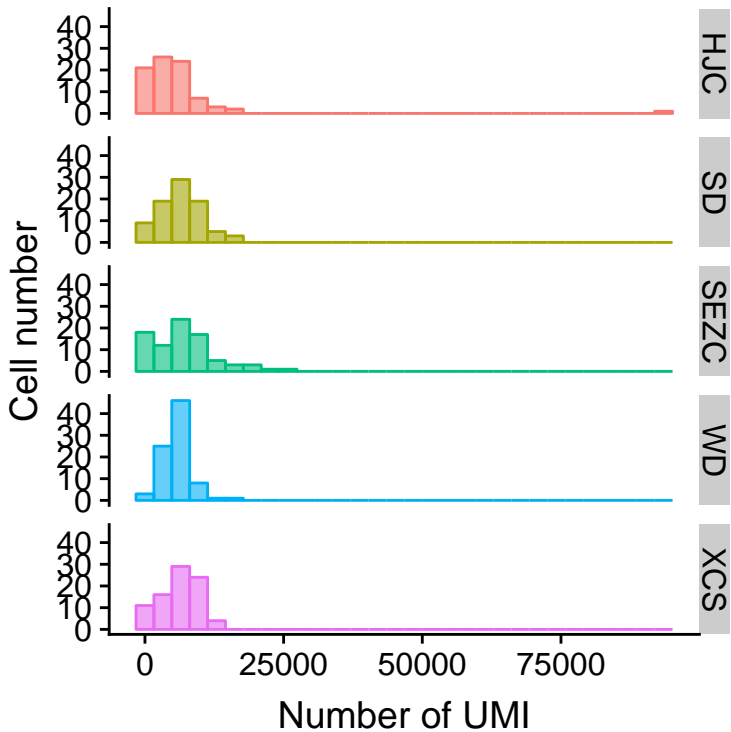
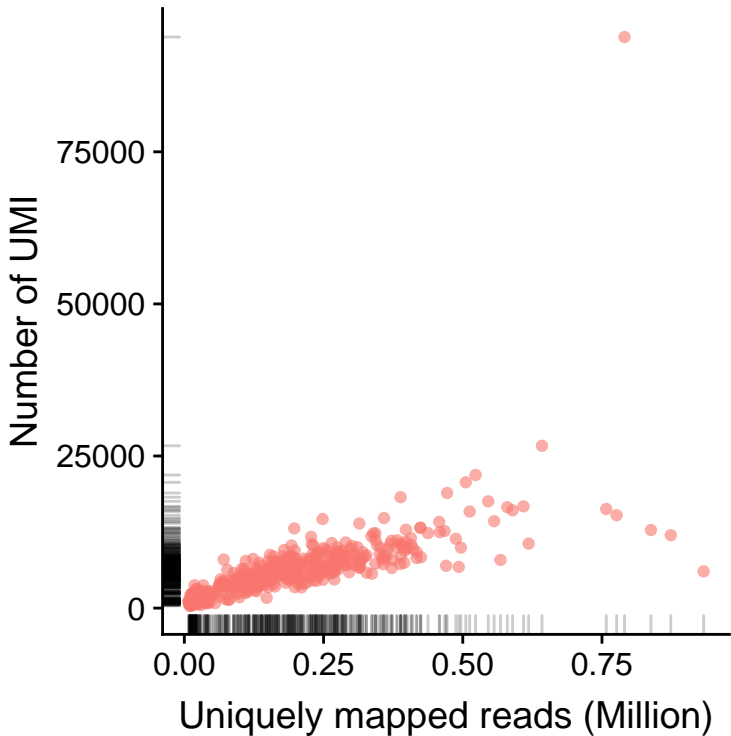


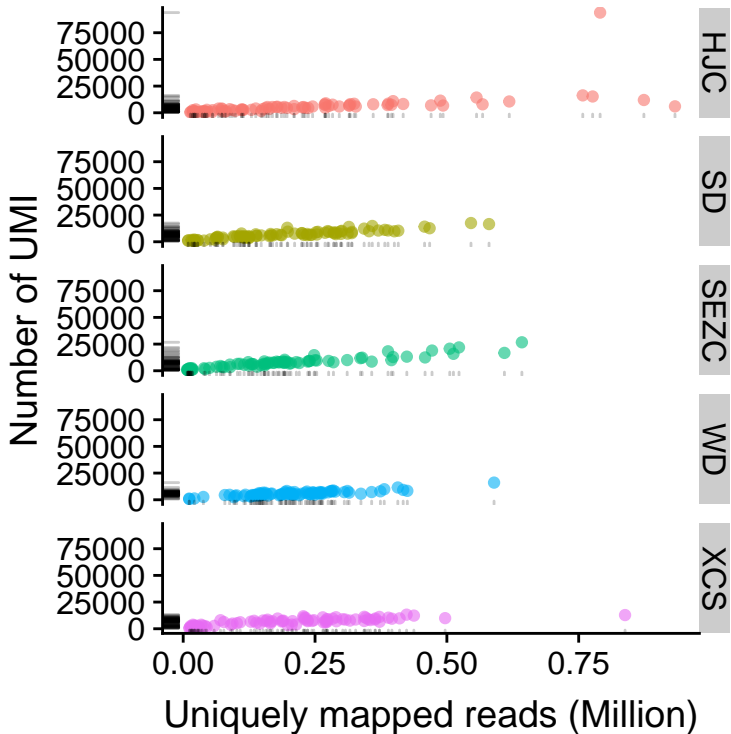
Number of detected genes

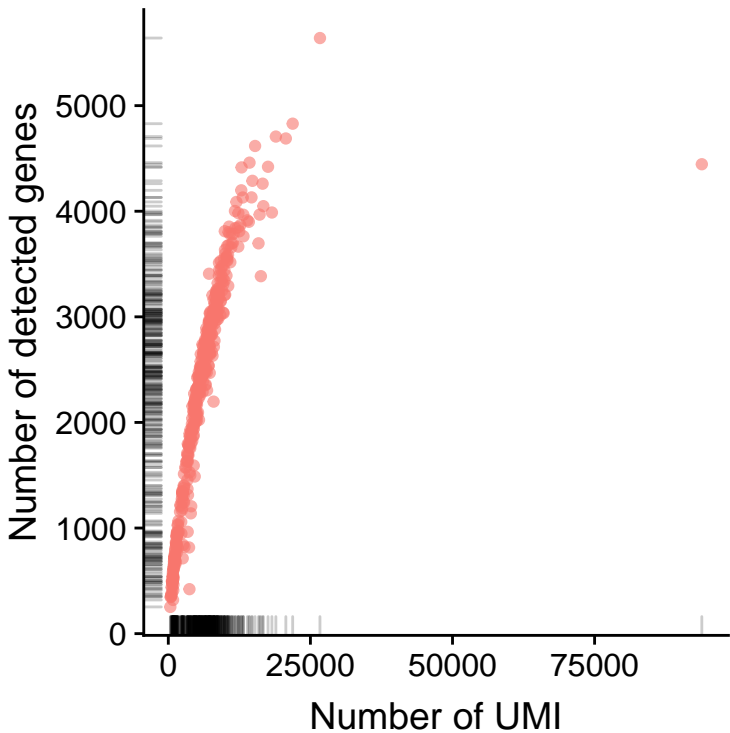




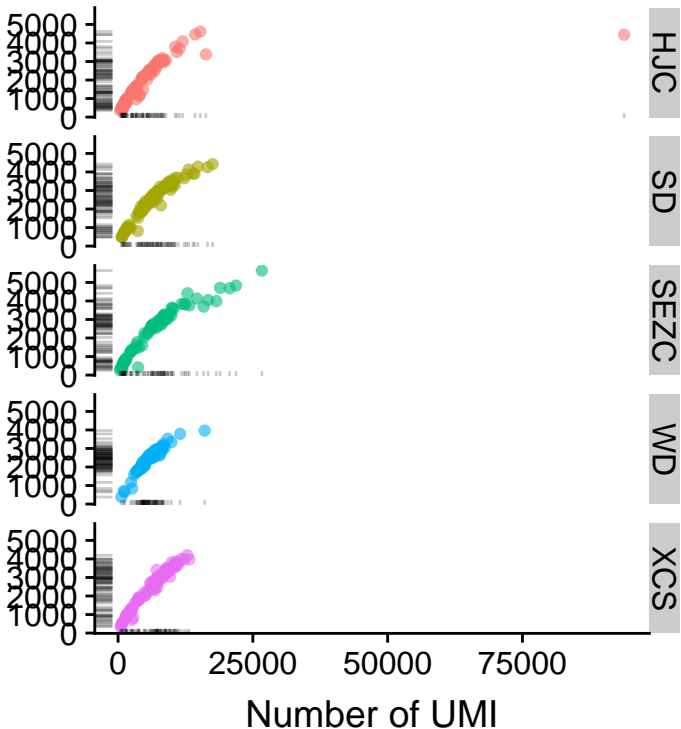


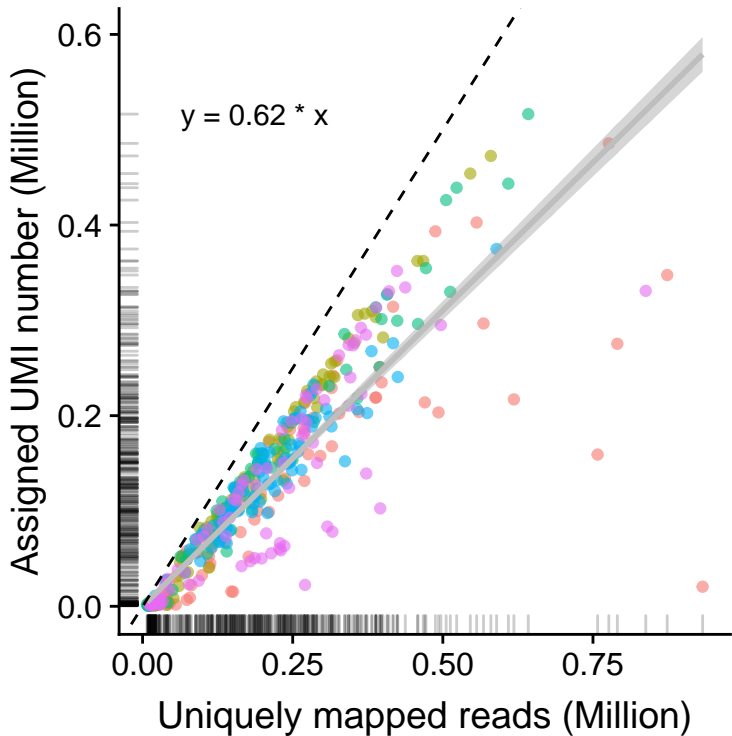


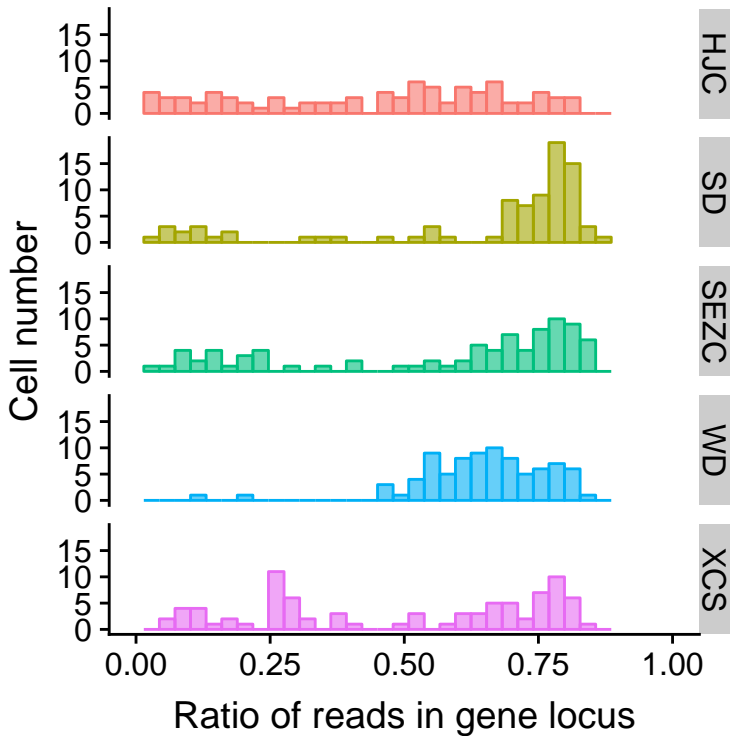




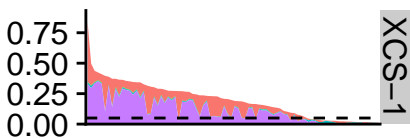
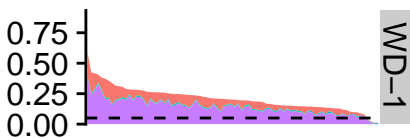
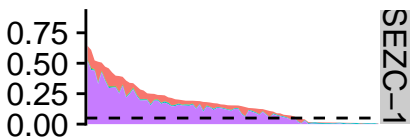
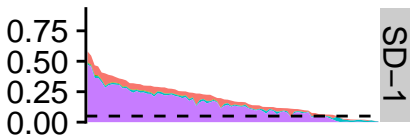
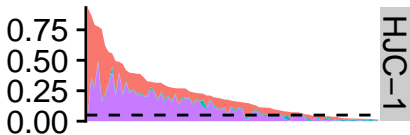
Number of detected genes



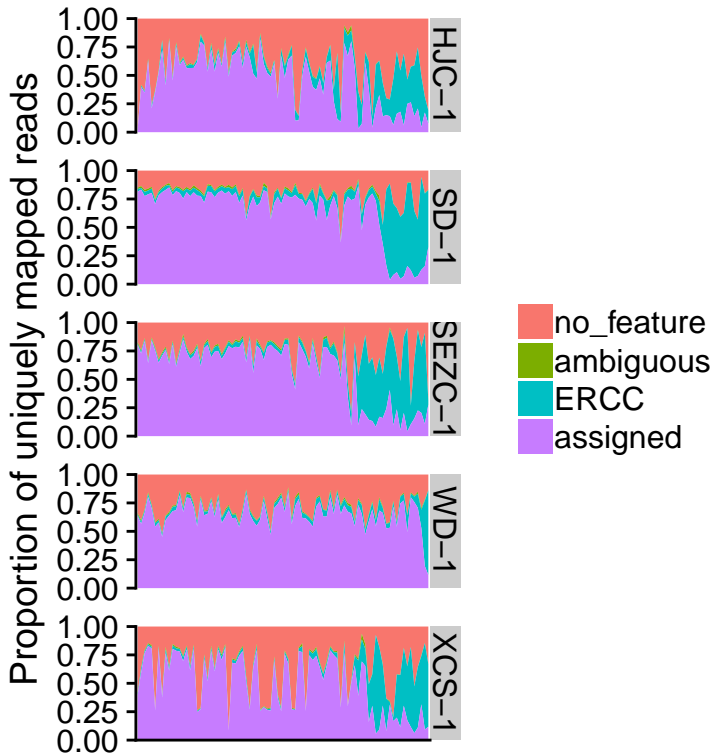


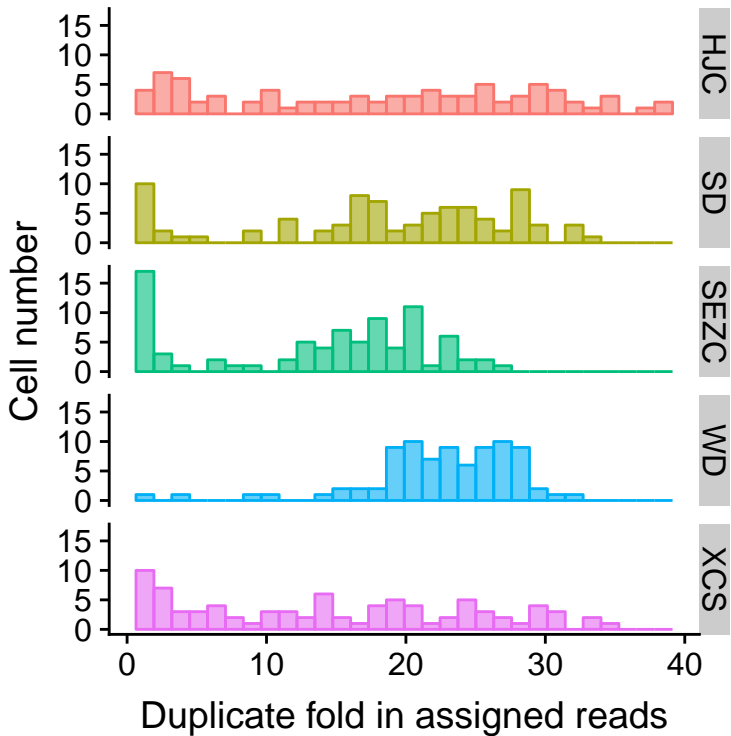


Uniquely mapped reads (Million)

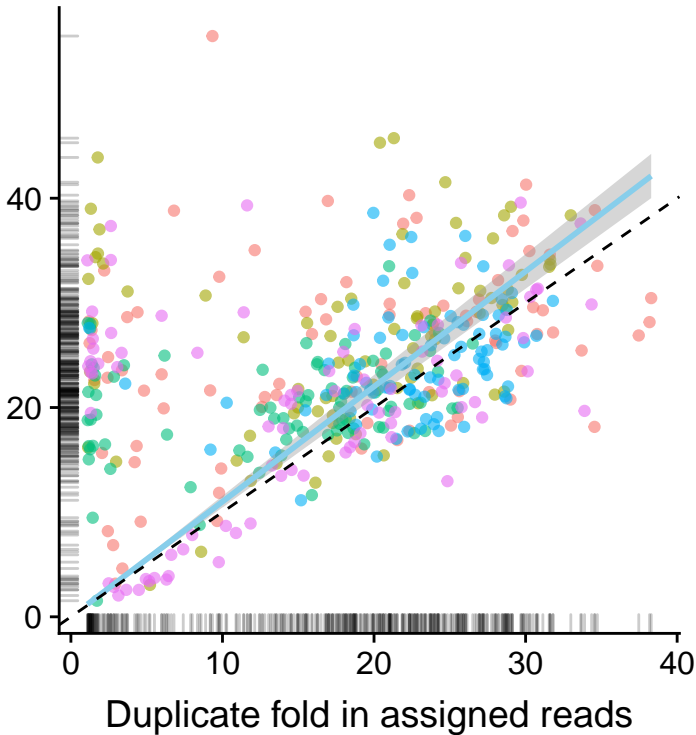


no_feature
ambiguous
ERCC
assigned

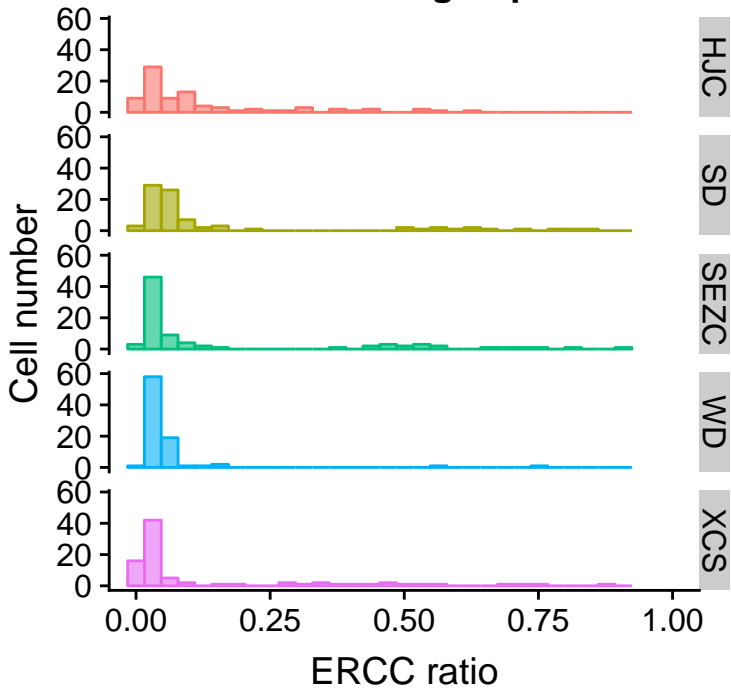




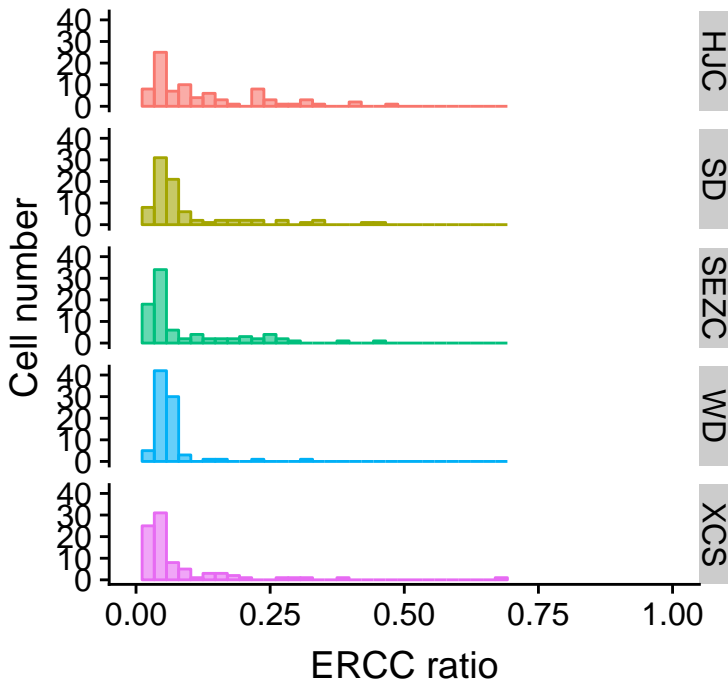
Duplicate fold in ERCC reads



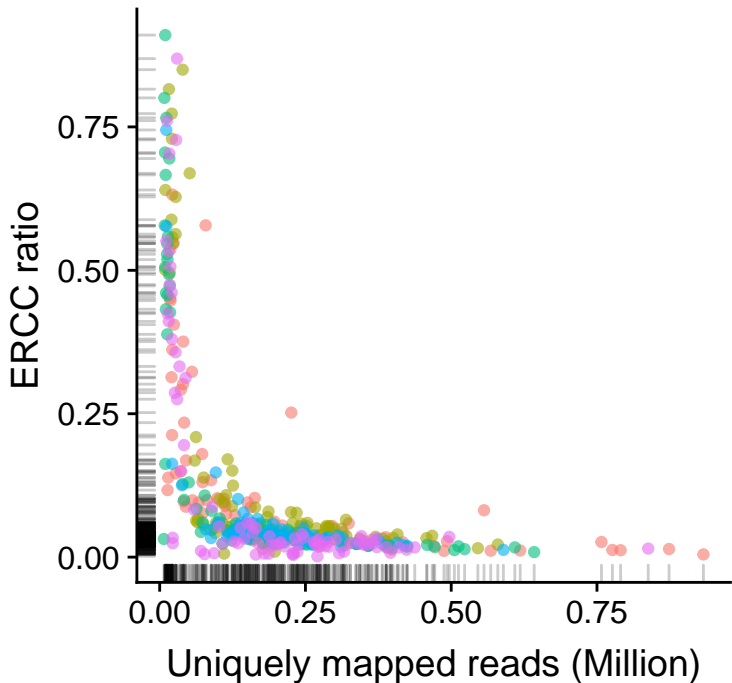
Before removing duplicates



After removing duplicates



Before removing duplicates



After removing duplicates

