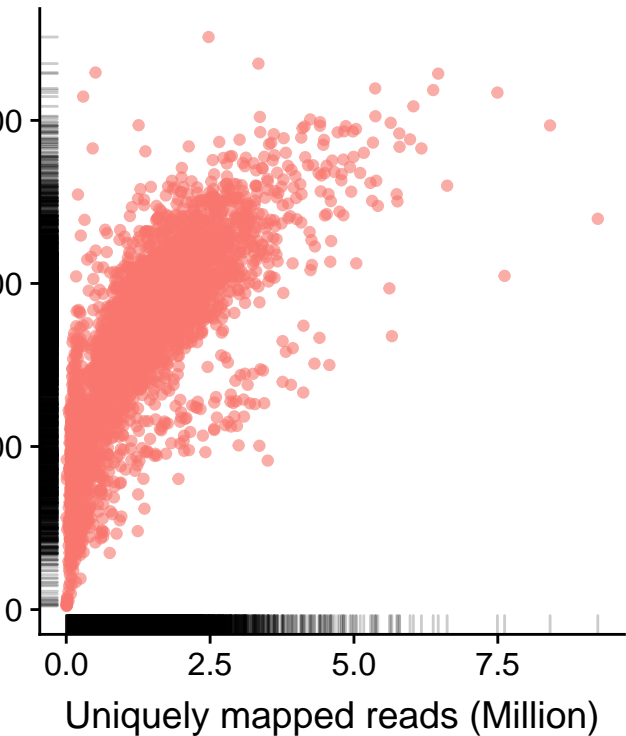
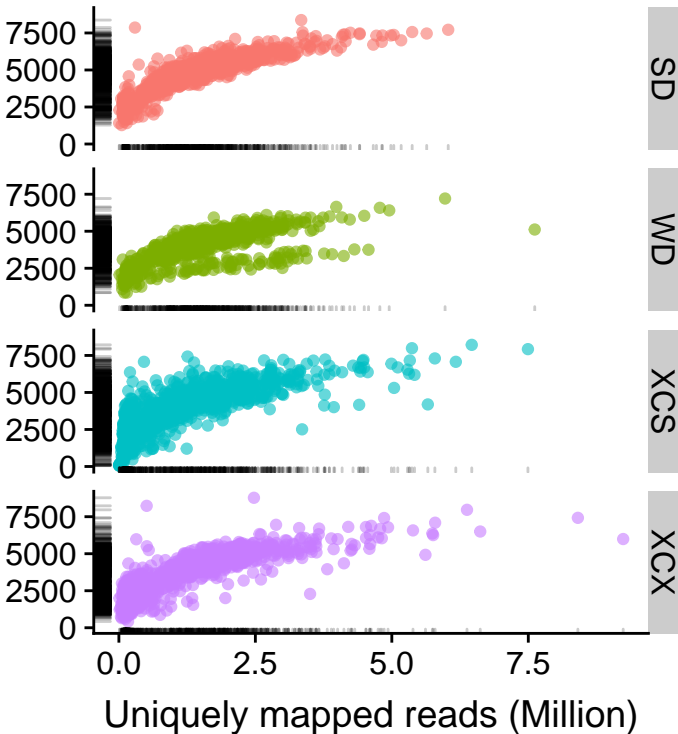
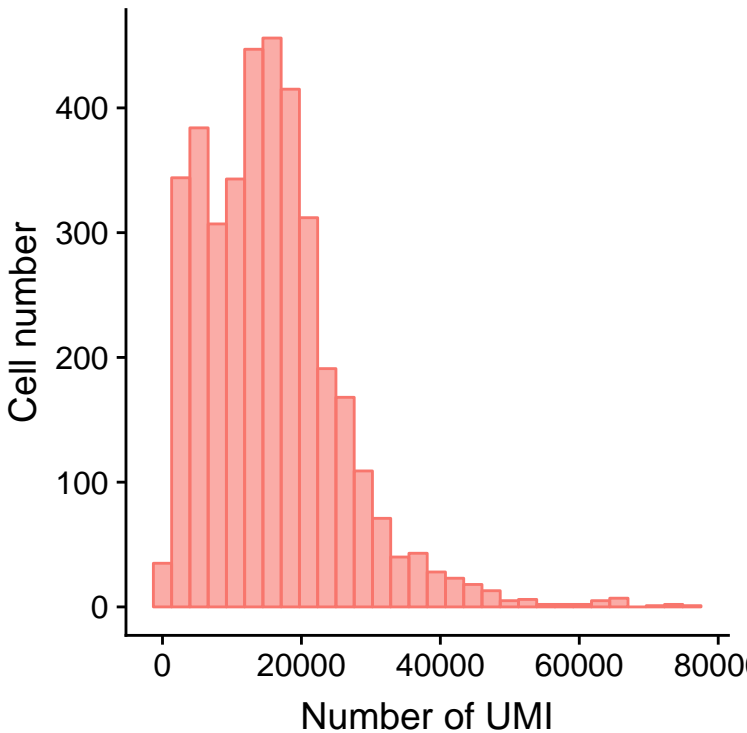


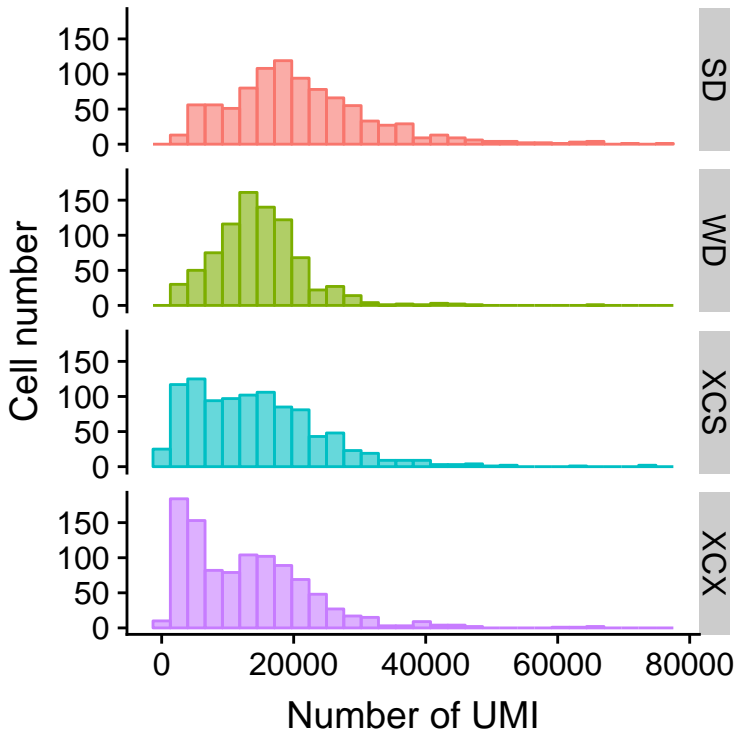
Number of detected genes

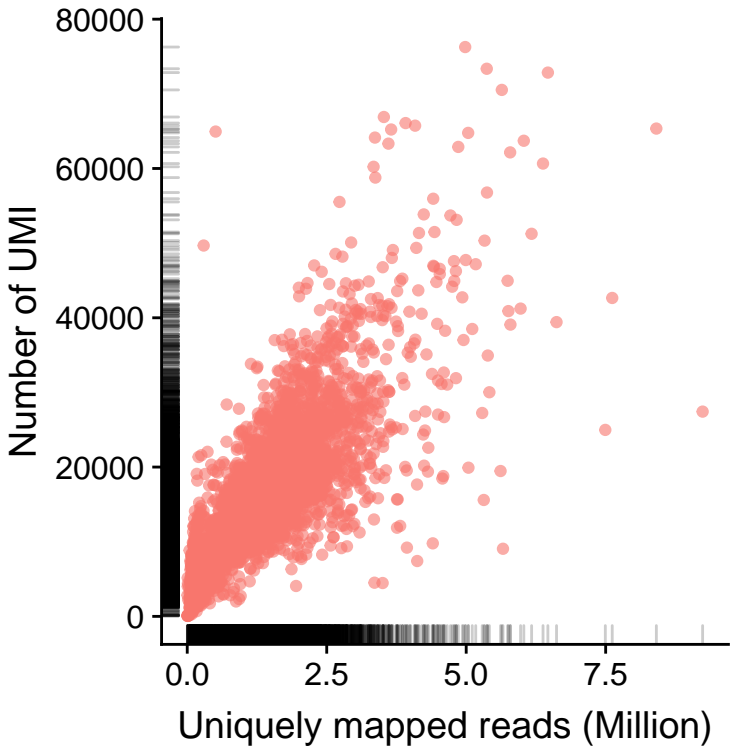


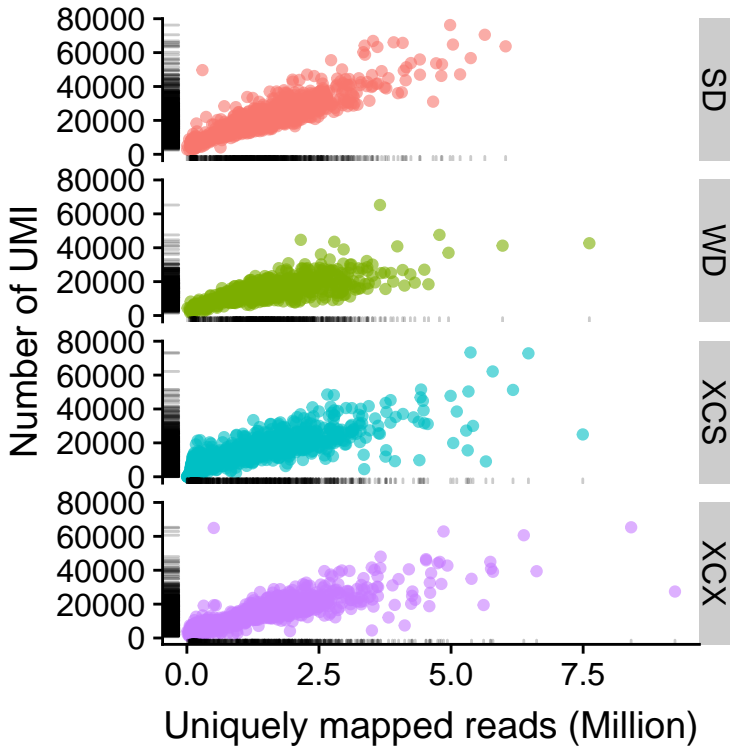
Number of detected genes



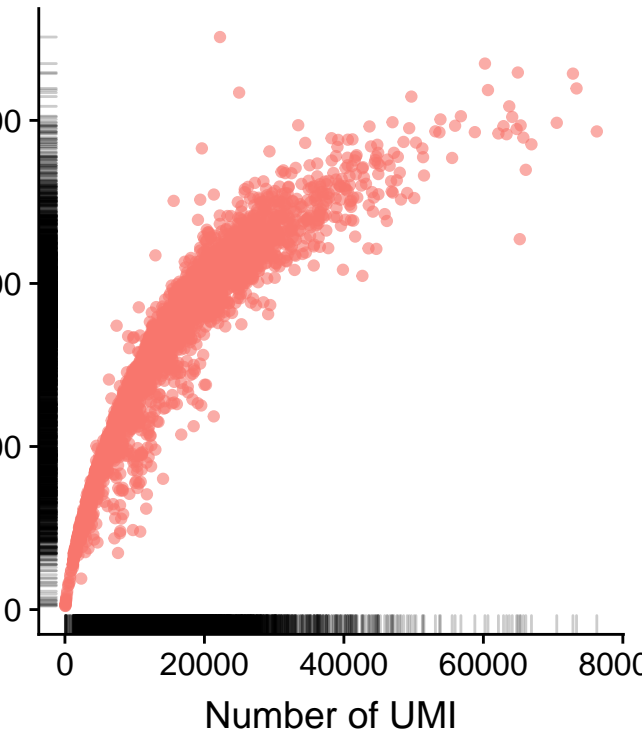




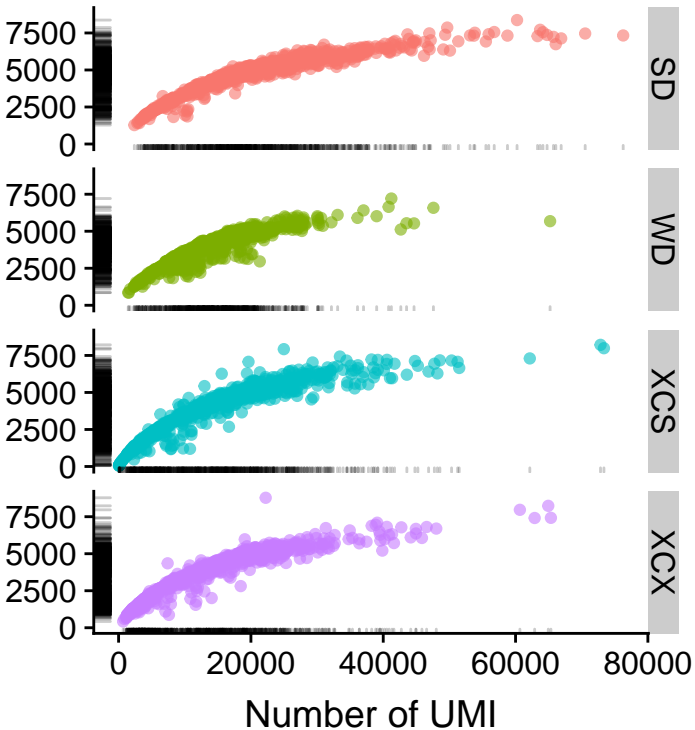


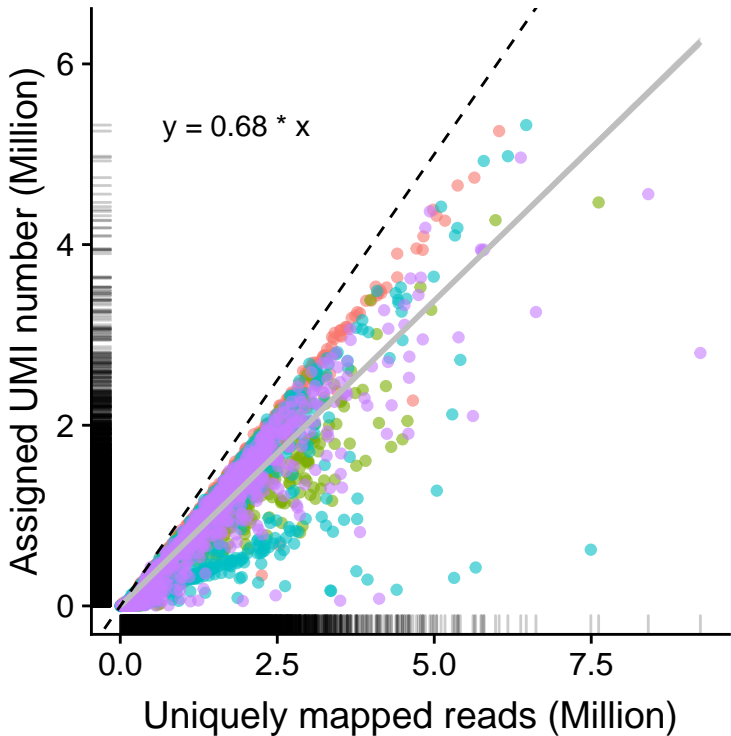


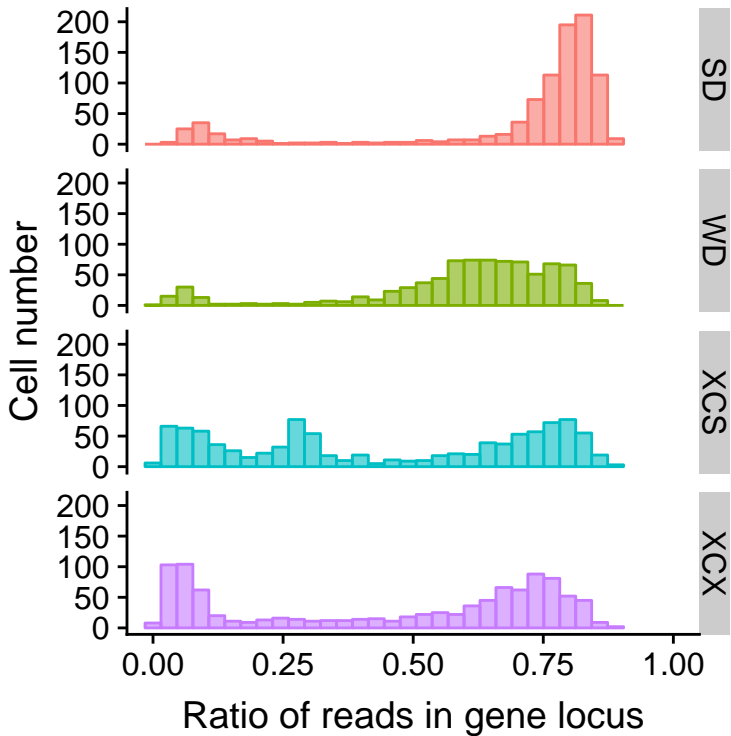
Number of detected genes



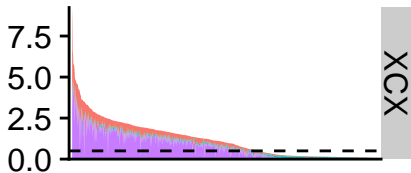
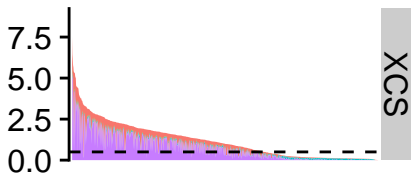
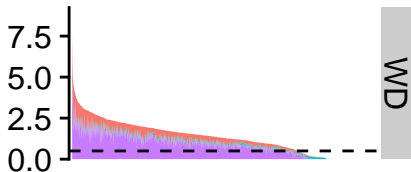
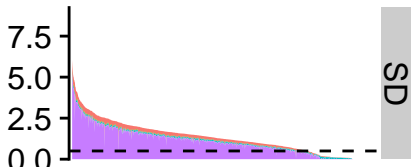
Number of detected genes



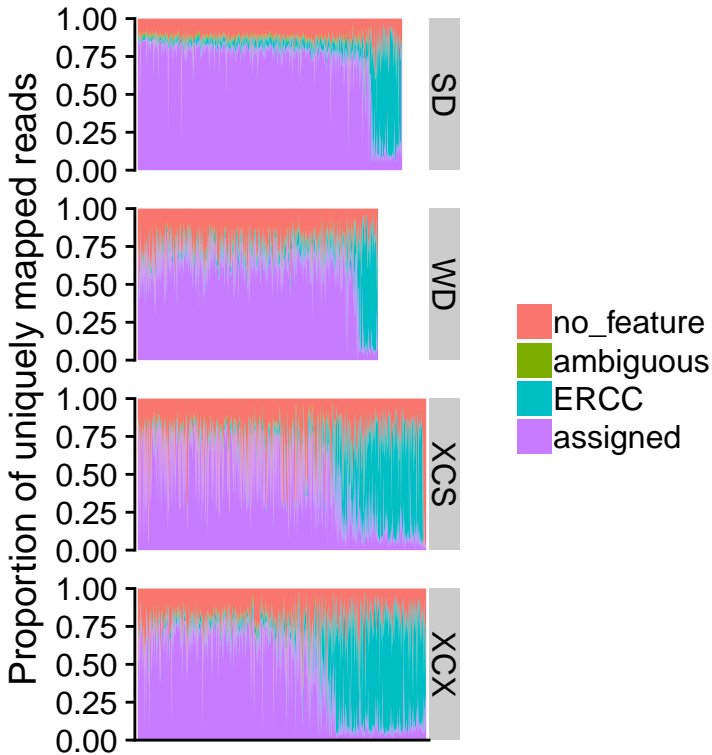


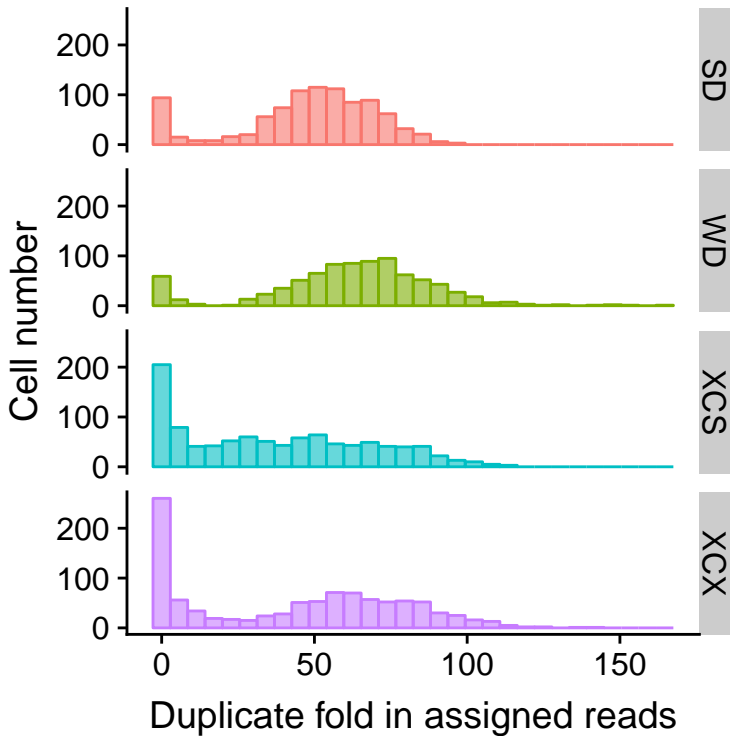


Uniquely mapped reads (Million)

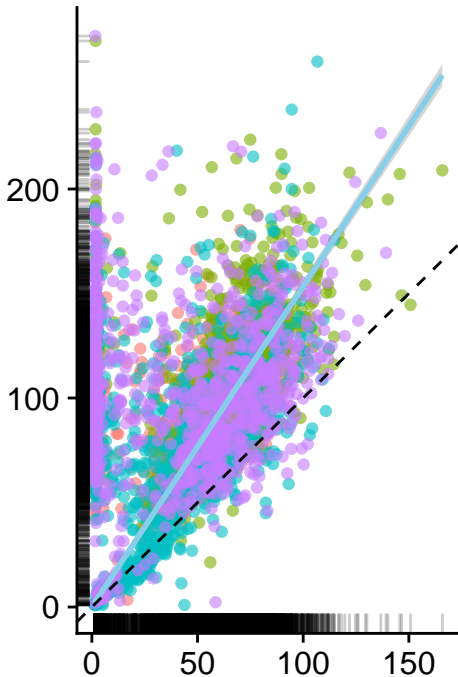


no_feature
ambiguous
ERCC
assigned





Duplicate fold in ERCC reads

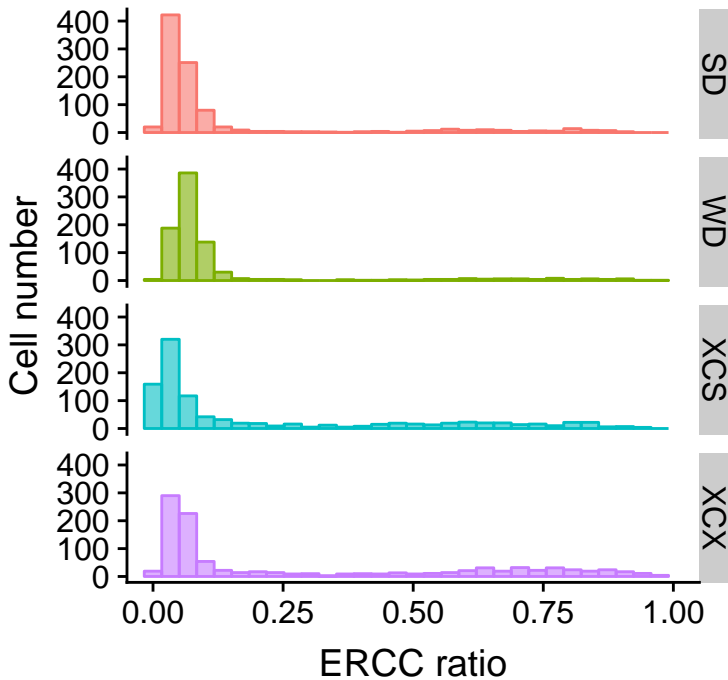


bigBatch

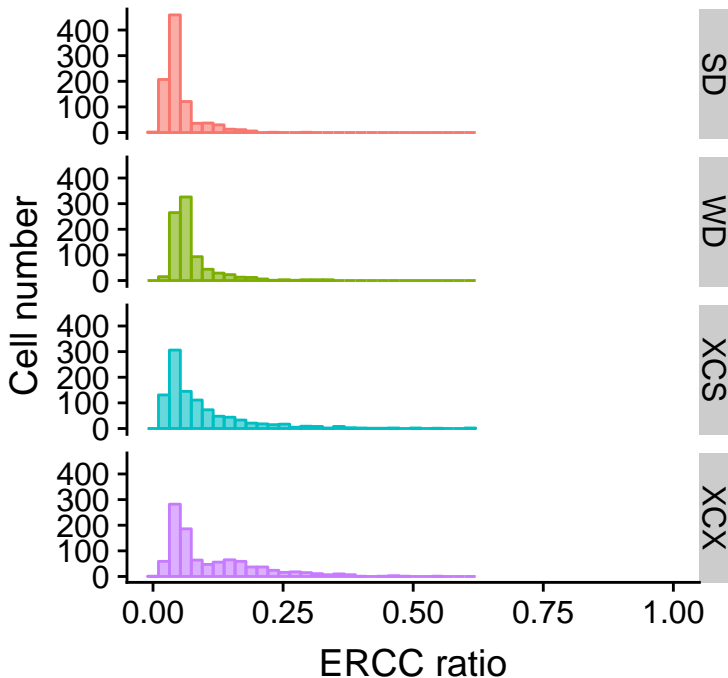
- SD
- WD
- XCS
- XCX

Duplicate fold in assigned reads

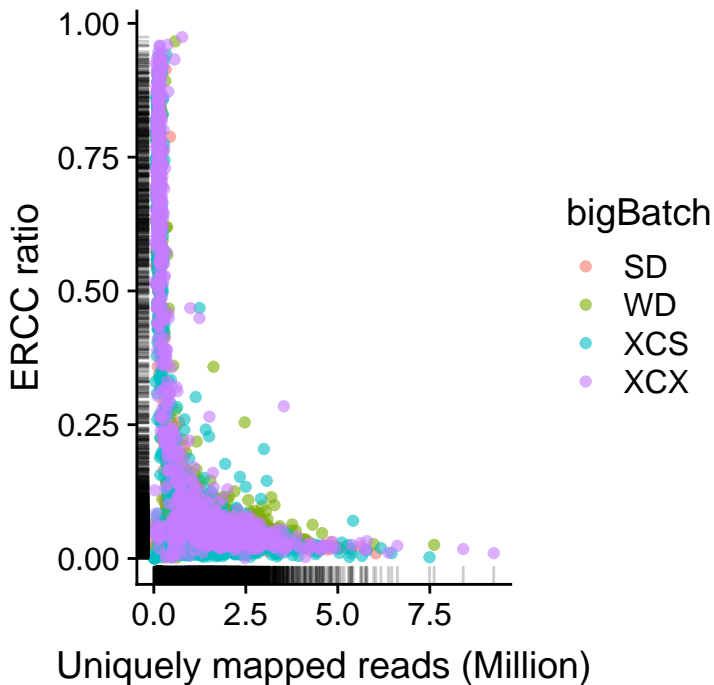
Before removing duplicates



After removing duplicates



Before removing duplicates



After removing duplicates

