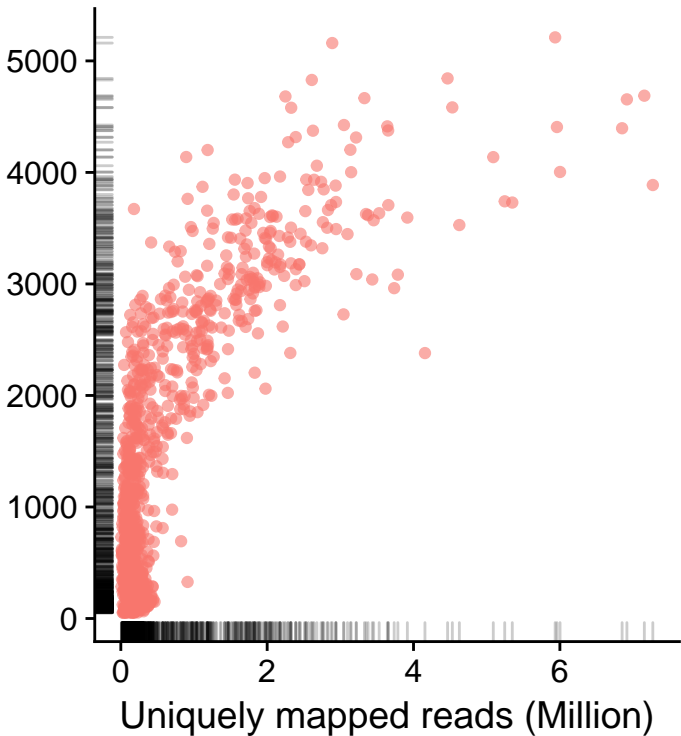
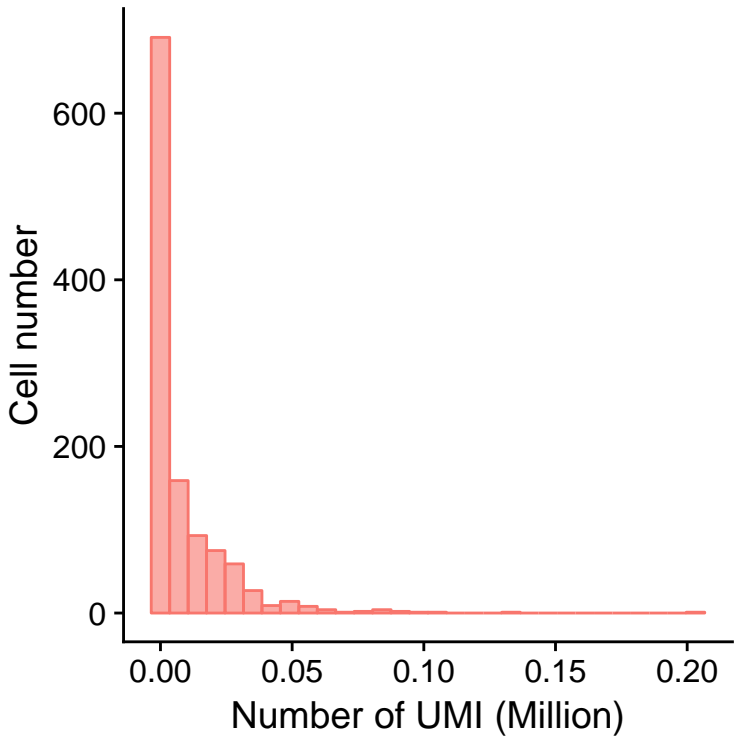
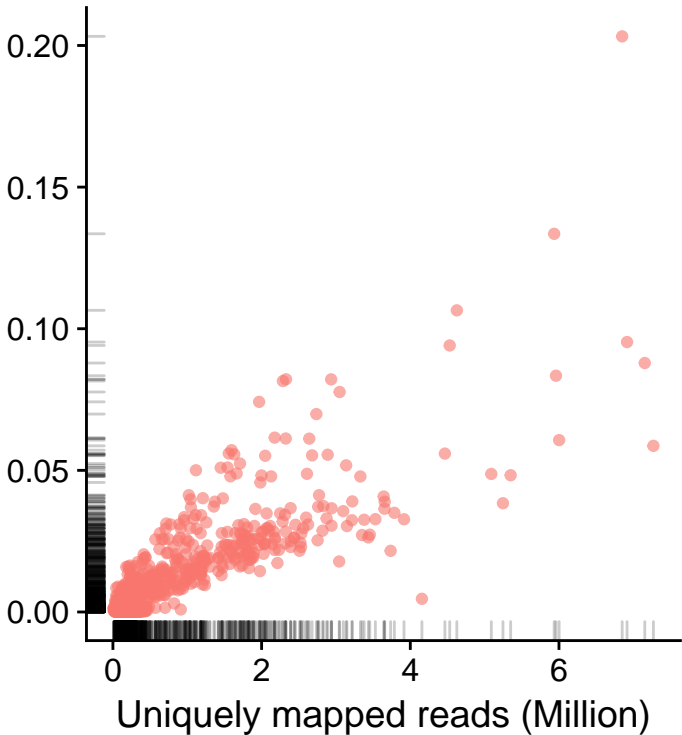


Number of detected genes

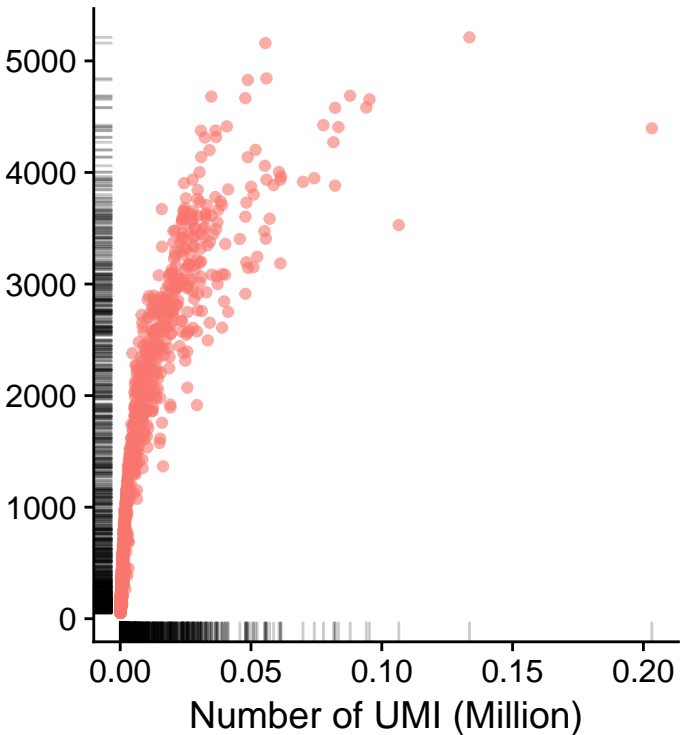


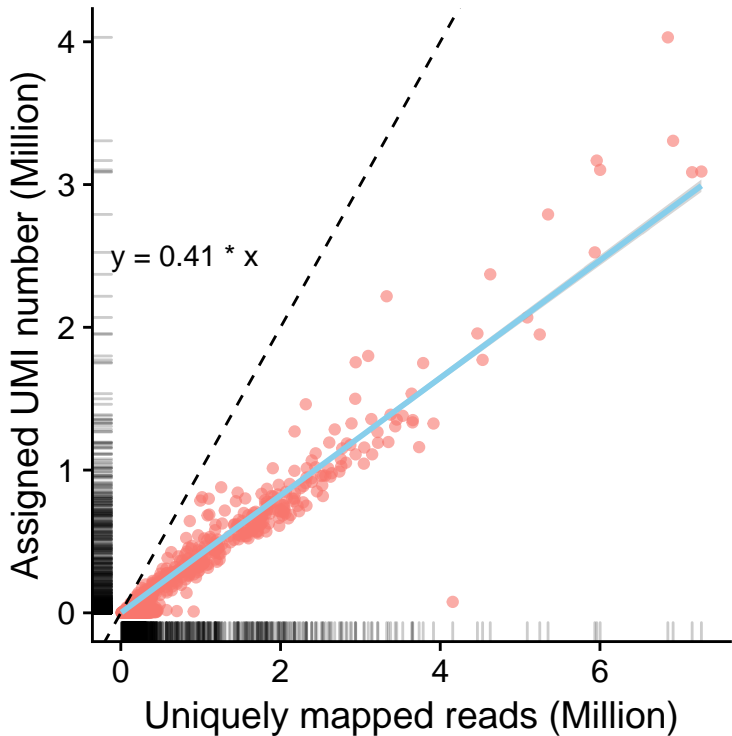


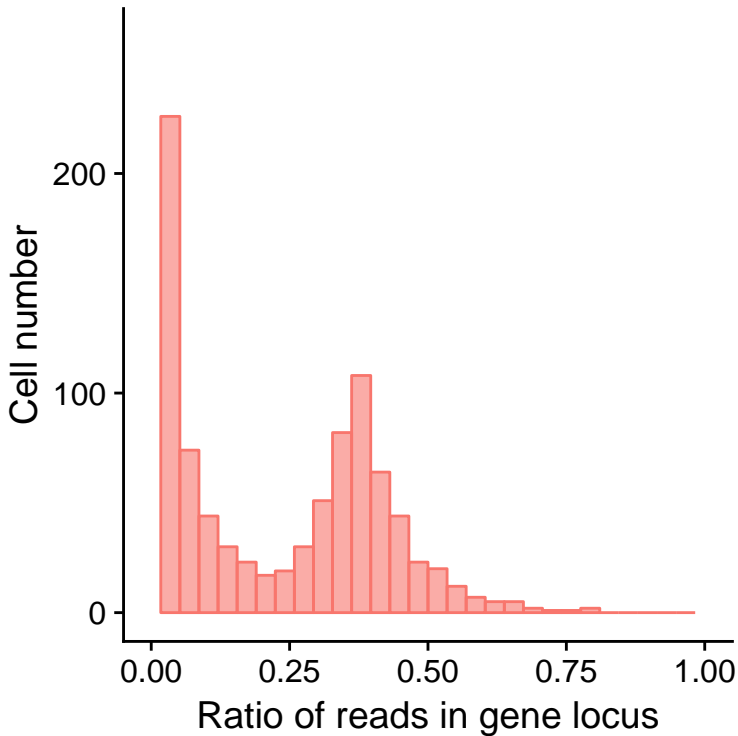
Number of UMI (Million)



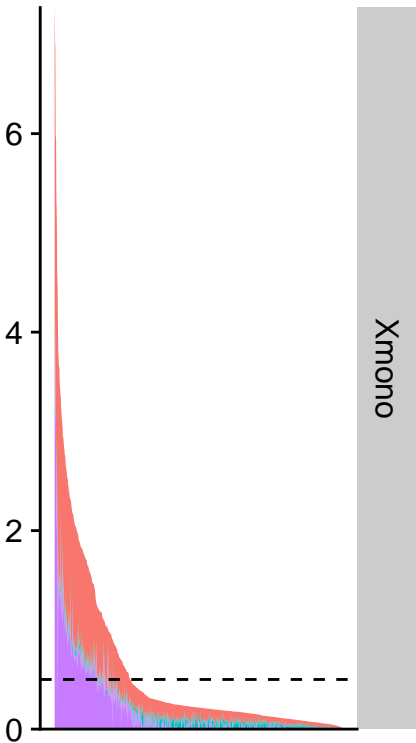
Number of detected genes





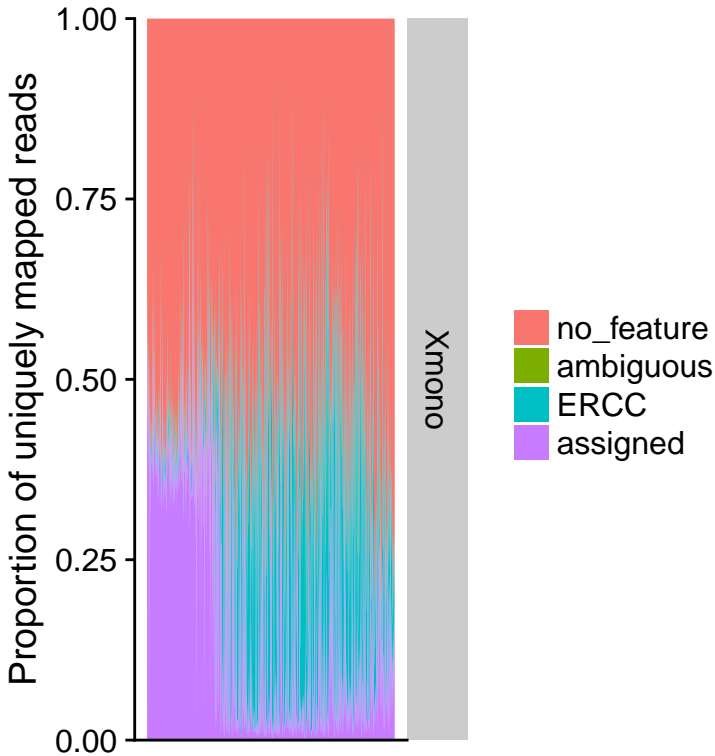


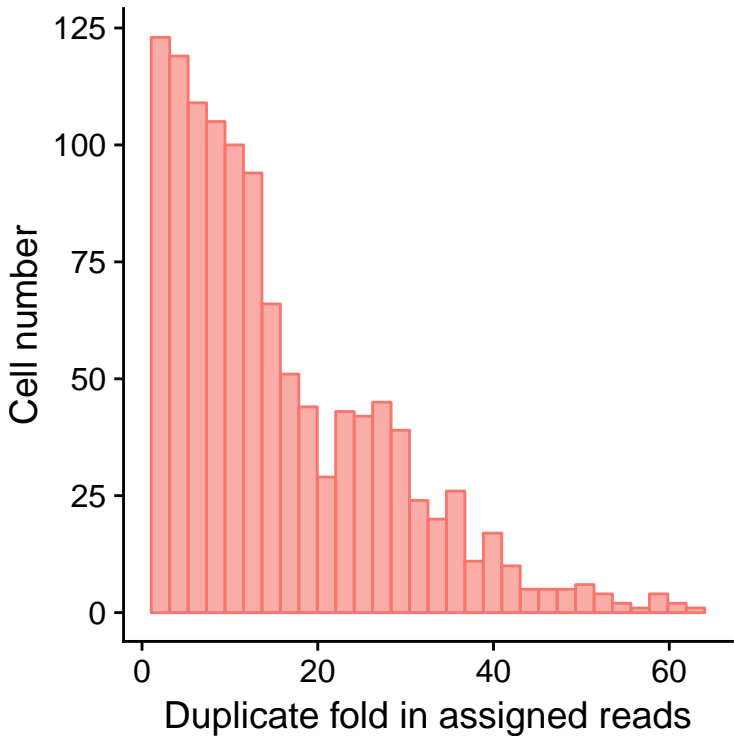
Uniquely mapped reads (Million)



Xmono

- no_feature
- ambiguous
- ERCC
- assigned





Duplicate fold in ERCC reads

75

50

25

0

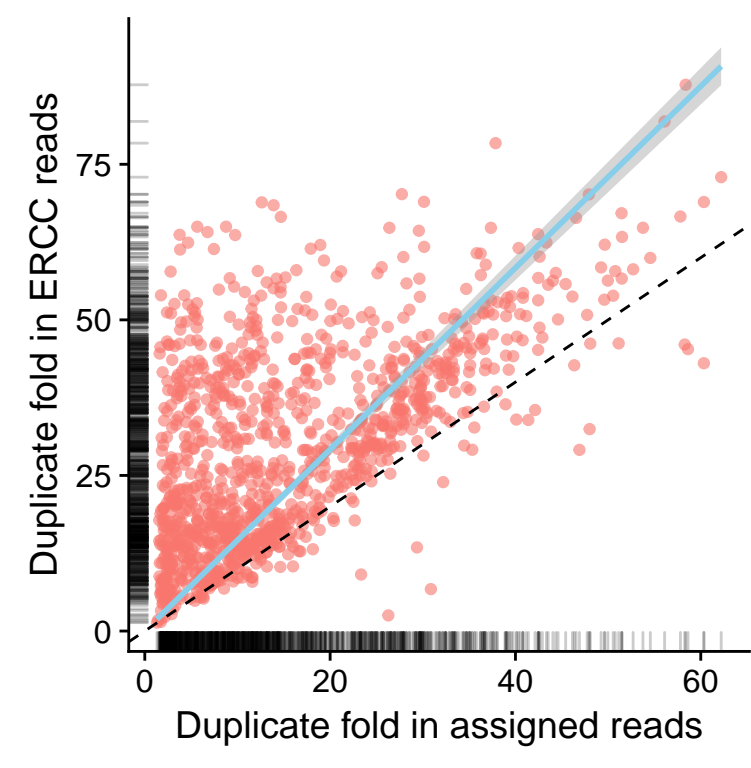
0

20

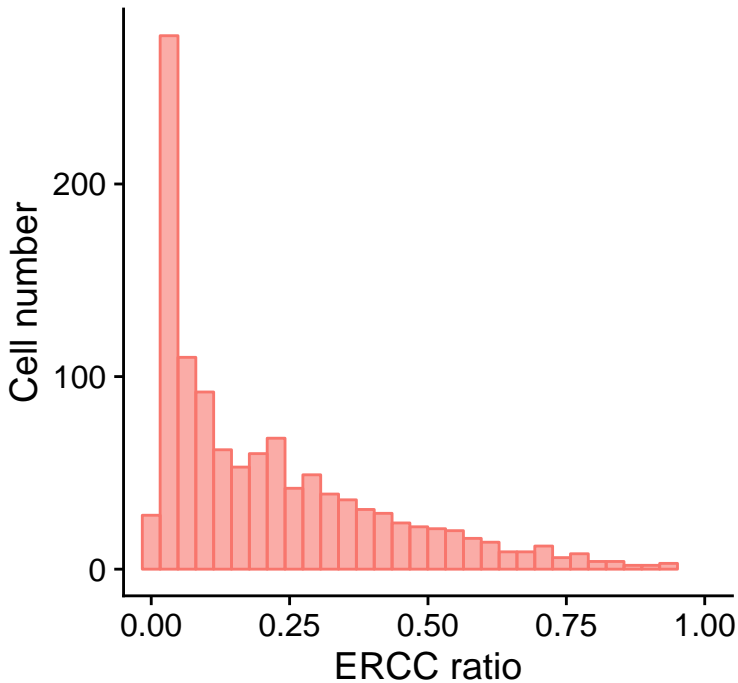
40

60

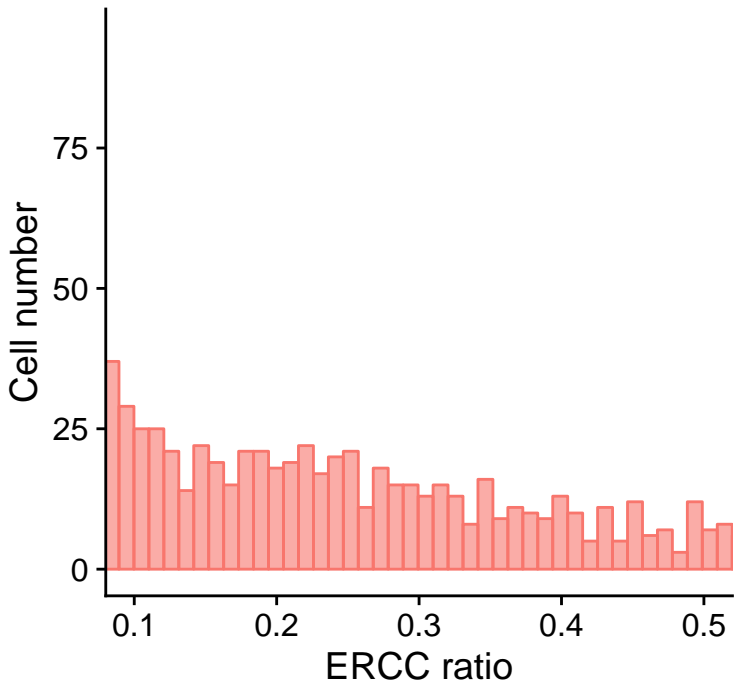
Duplicate fold in assigned reads



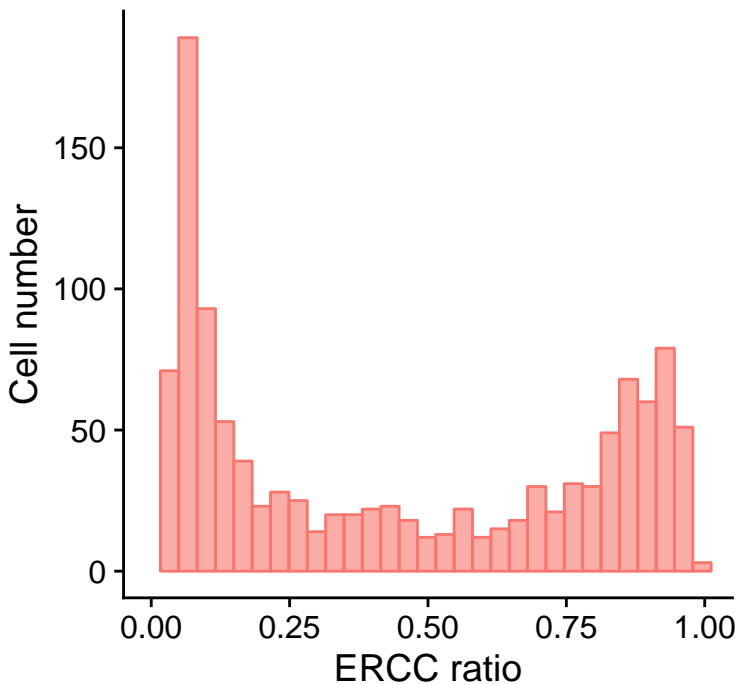
Before removing duplicates



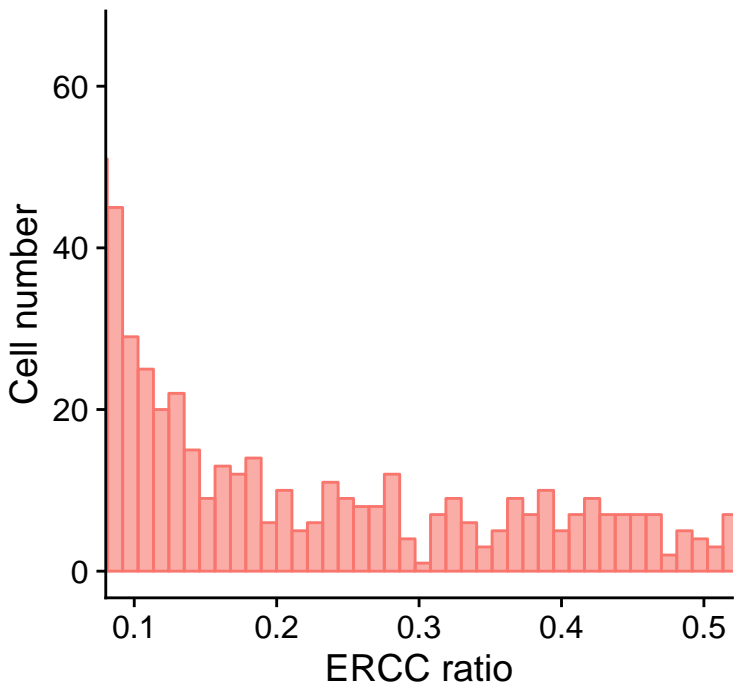
Before removing dup. (Zoom in)



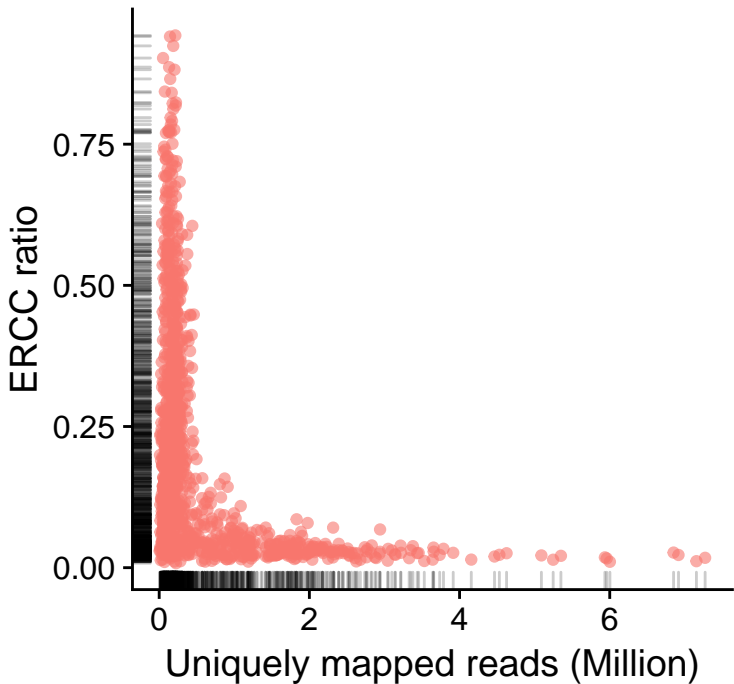
After removing duplicates



After removing dup. (Zoom in)



Before removing duplicates



After removing duplicates

