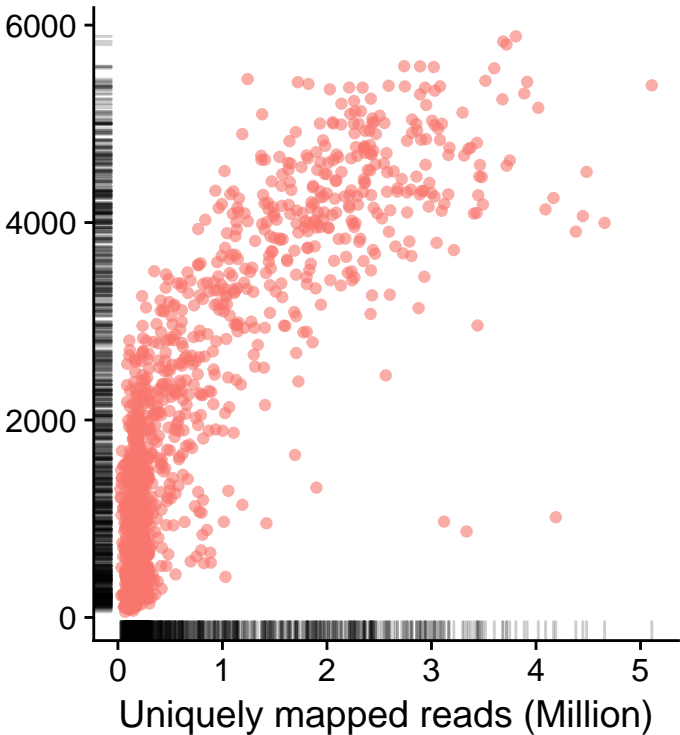
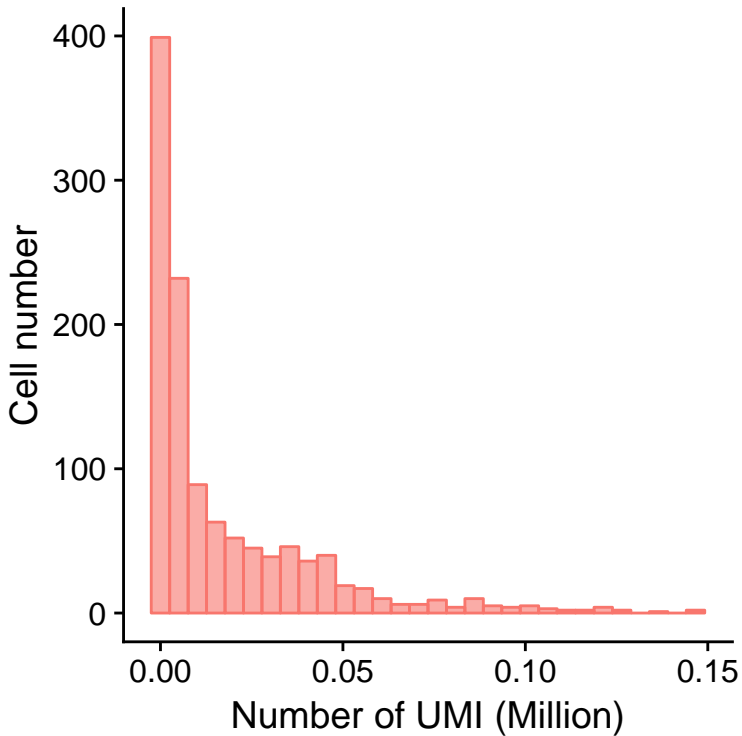
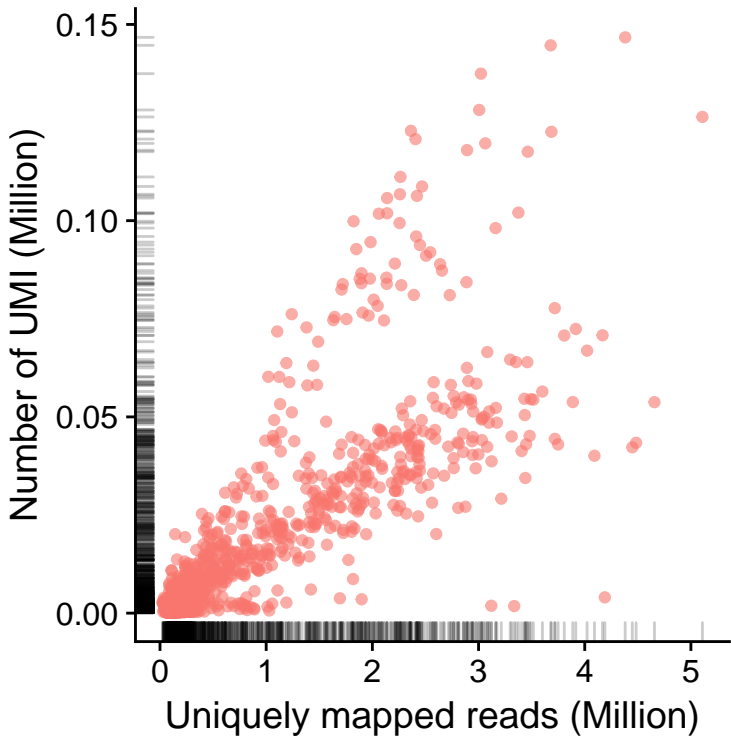


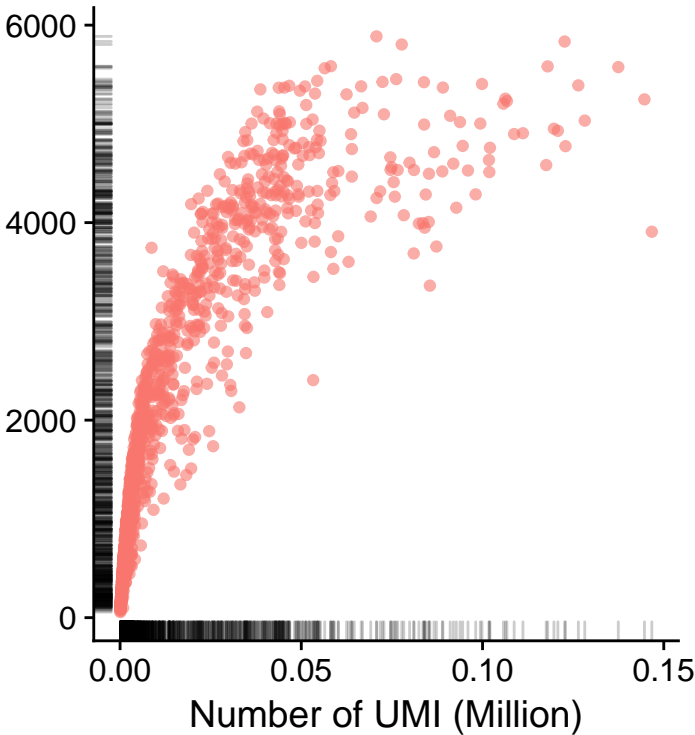
Number of detected genes

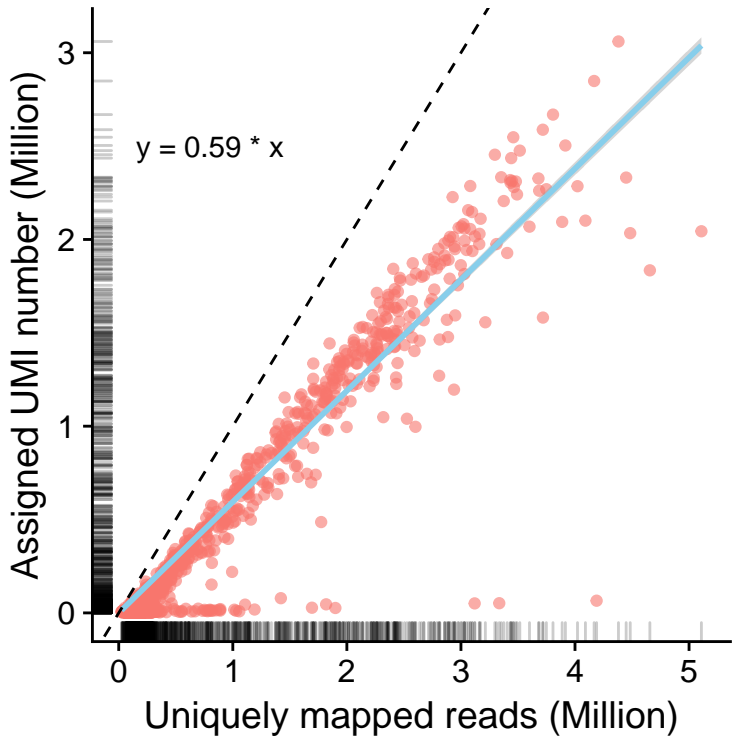


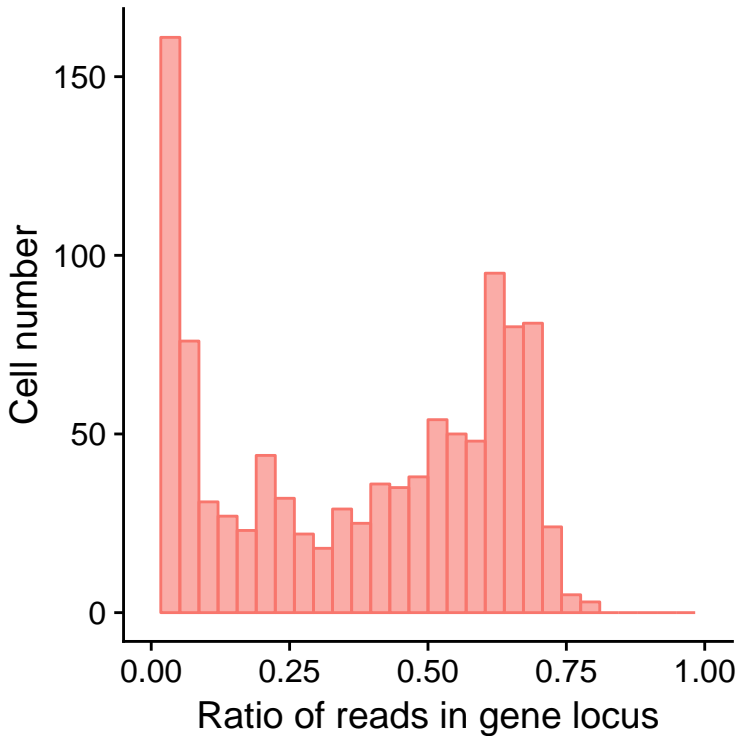




Number of detected genes





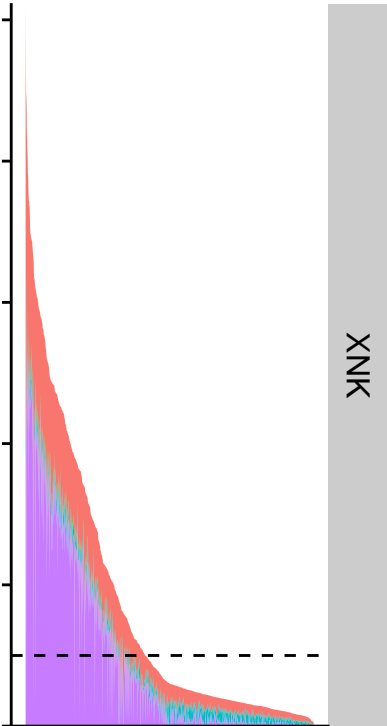


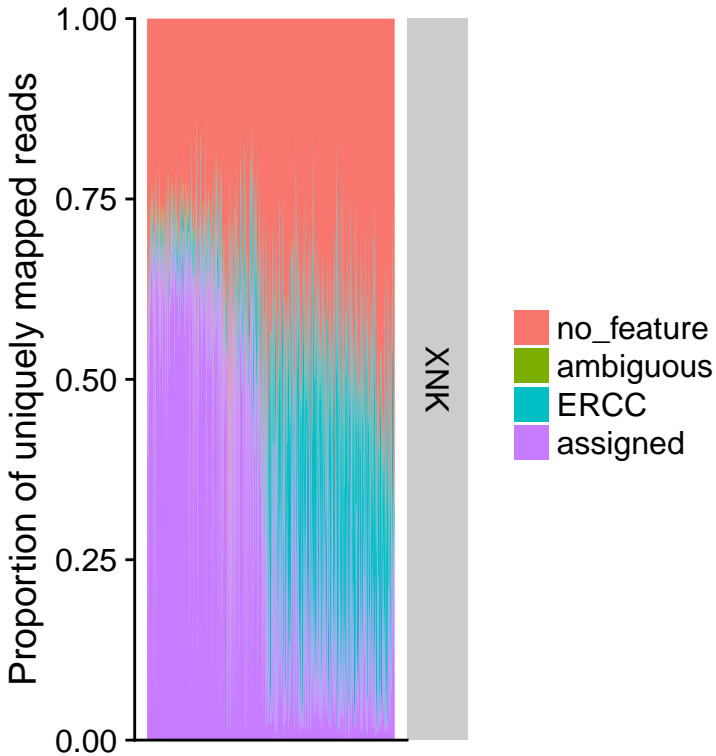
Uniquely mapped reads (Million)

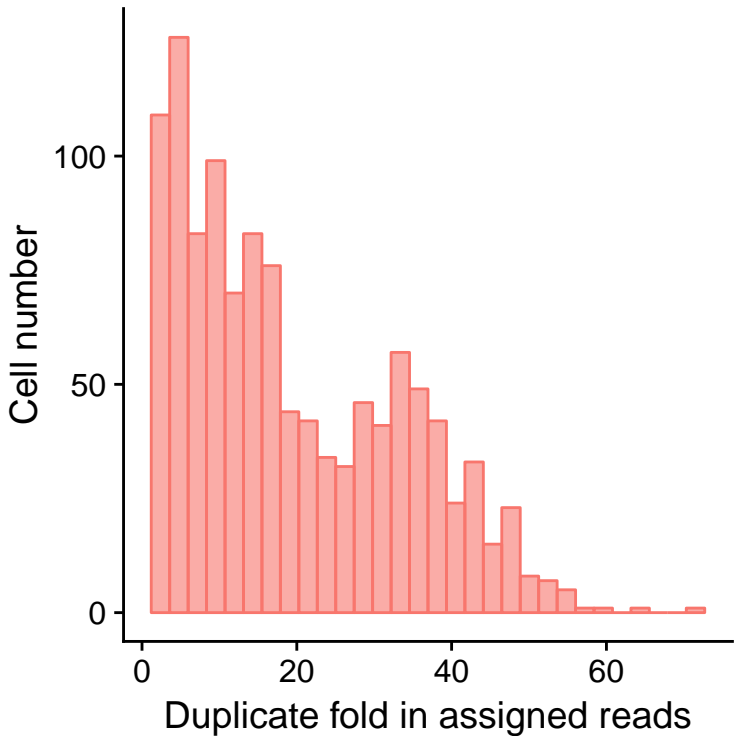
5
4
3
2
1
0

XNK

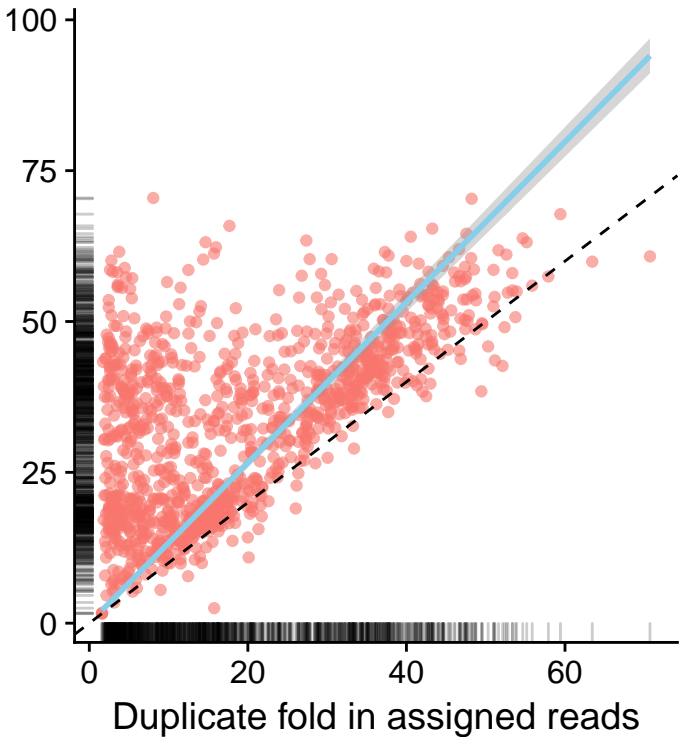
- no_feature
- ambiguous
- ERCC
- assigned



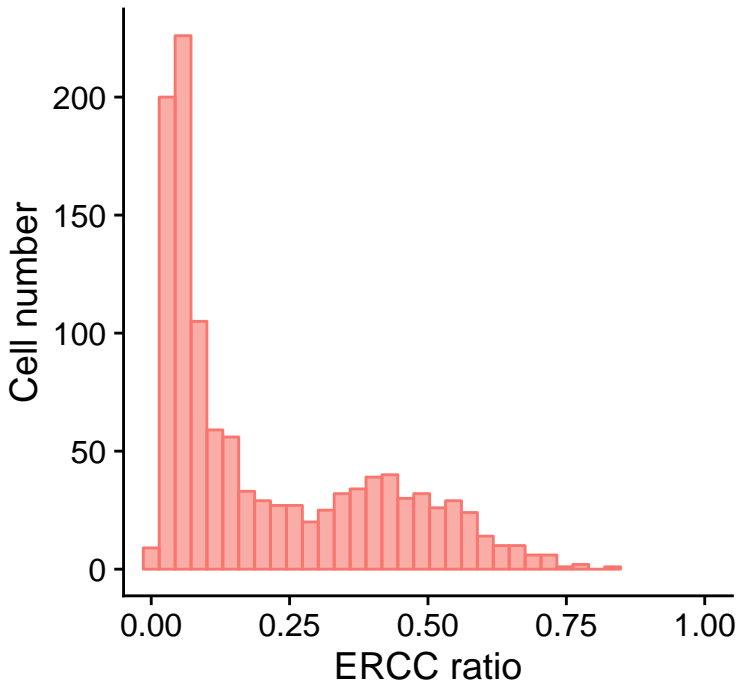




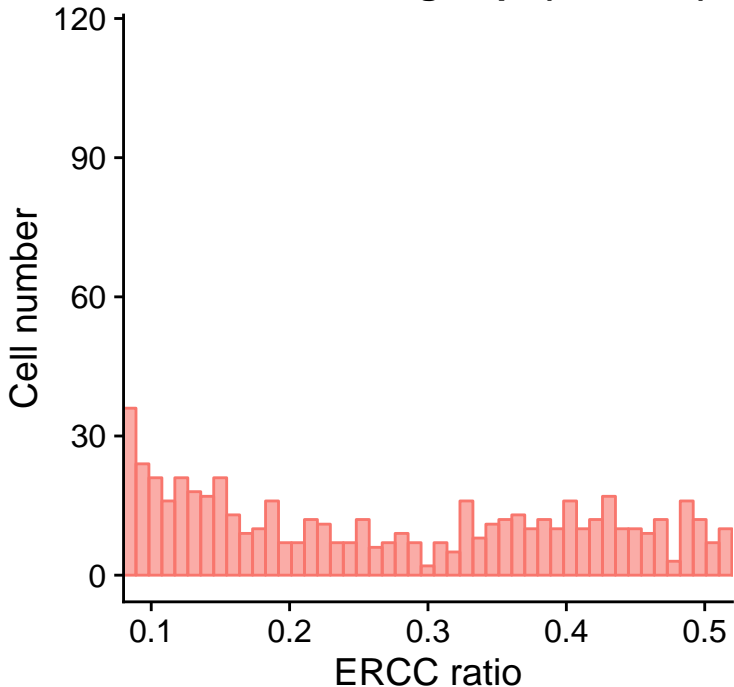
Duplicate fold in ERCC reads



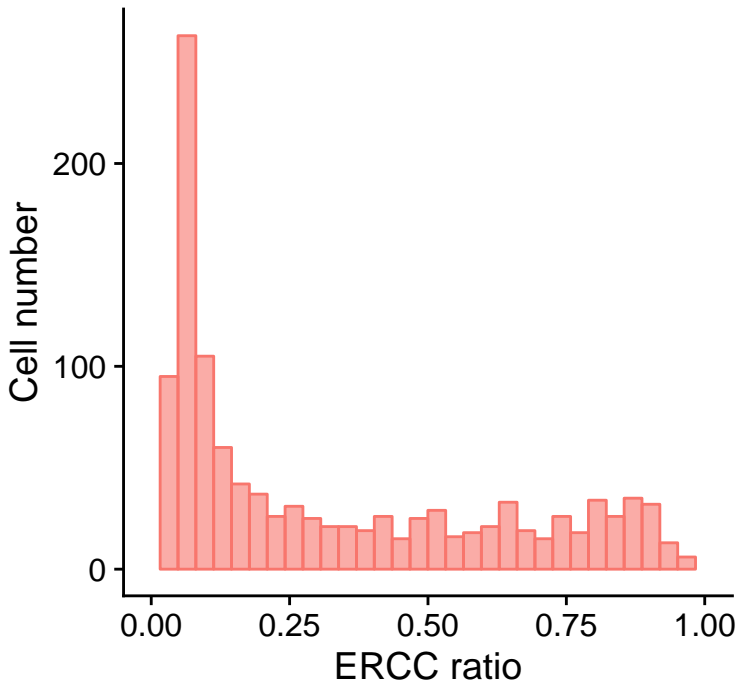
Before removing duplicates



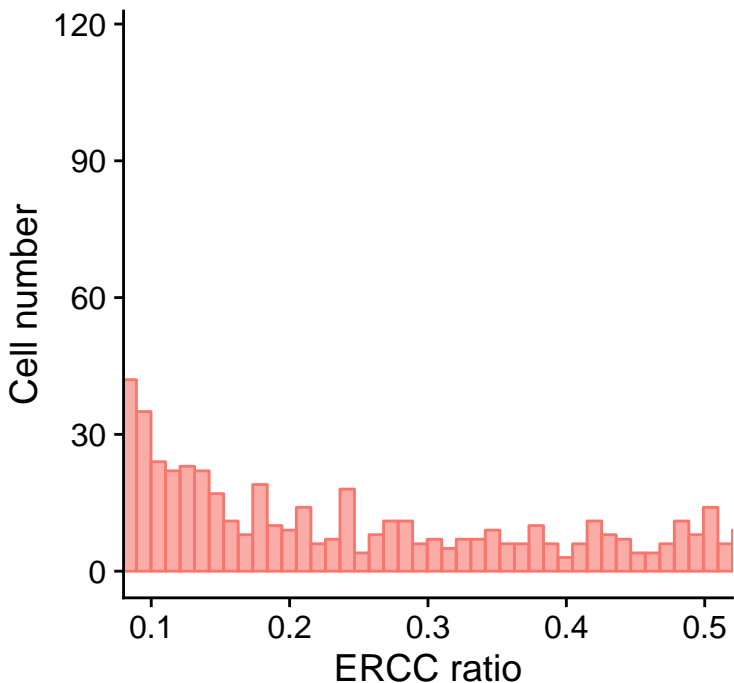
Before removing dup. (Zoom in)



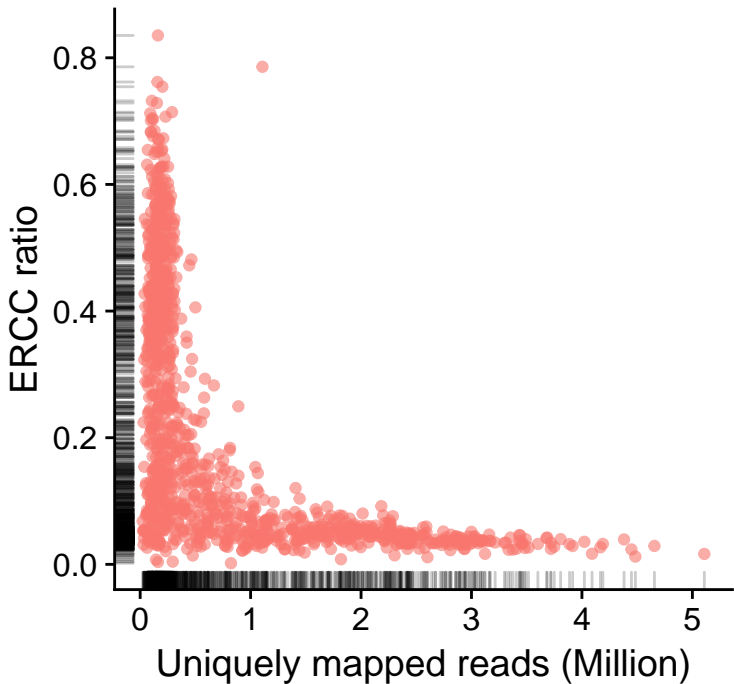
After removing duplicates



After removing dup. (Zoom in)



Before removing duplicates



After removing duplicates

