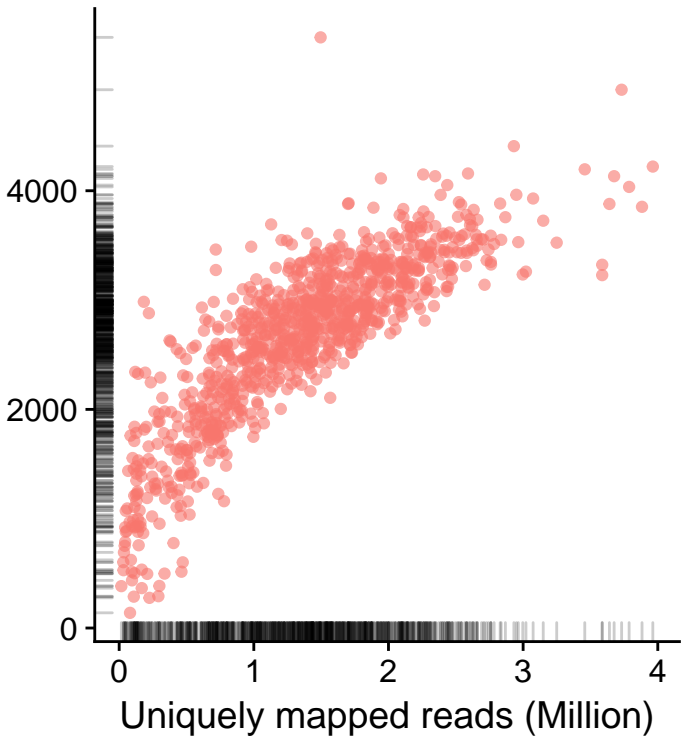
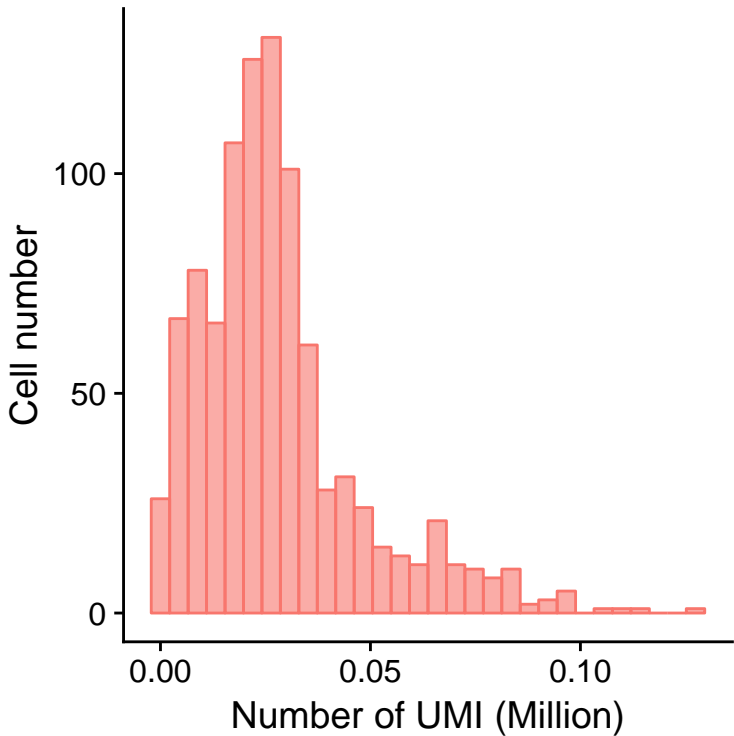


Number of detected genes





Number of UMI (Million)

0.10

0.05

0.00

0

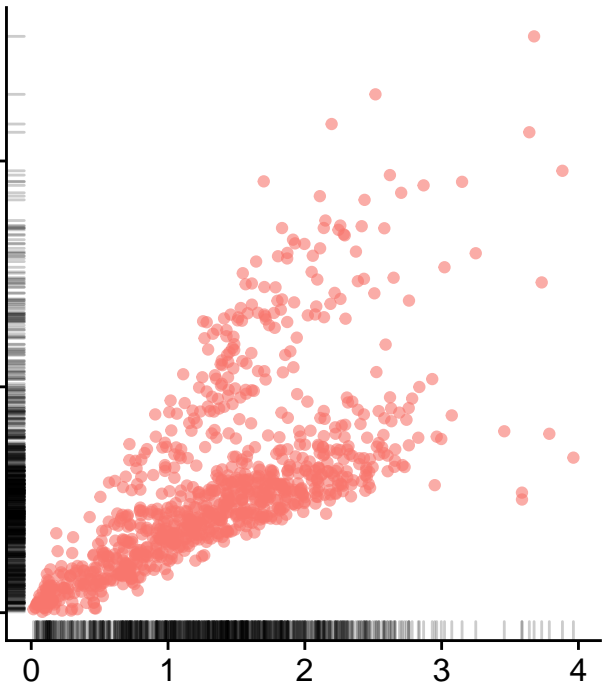
1

2

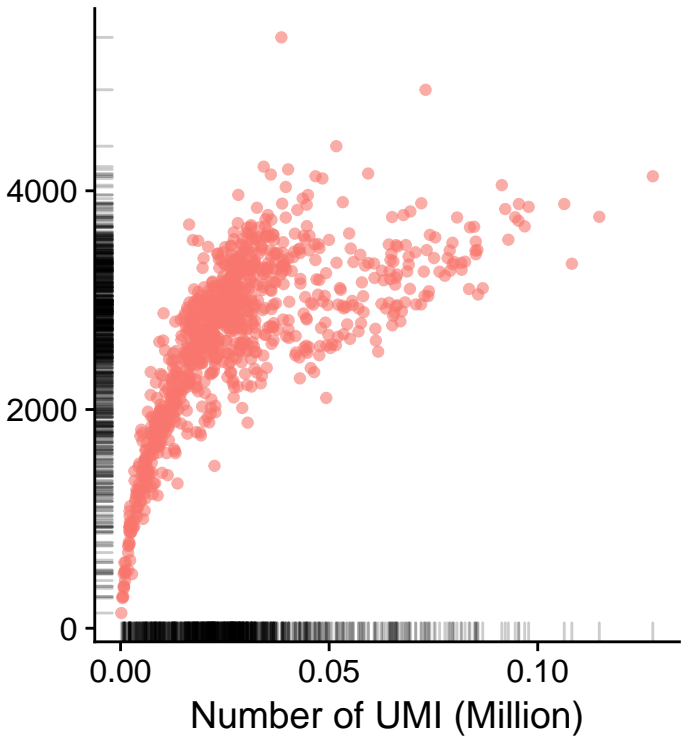
3

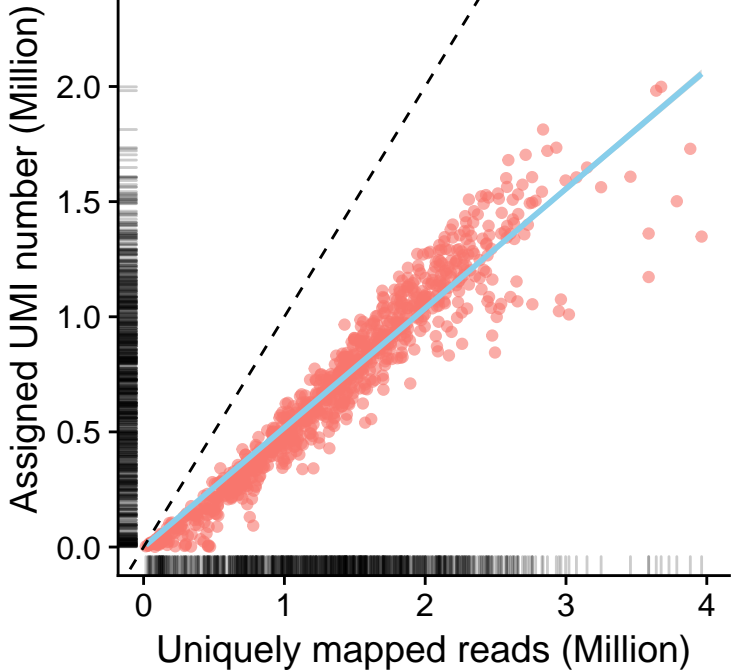
4

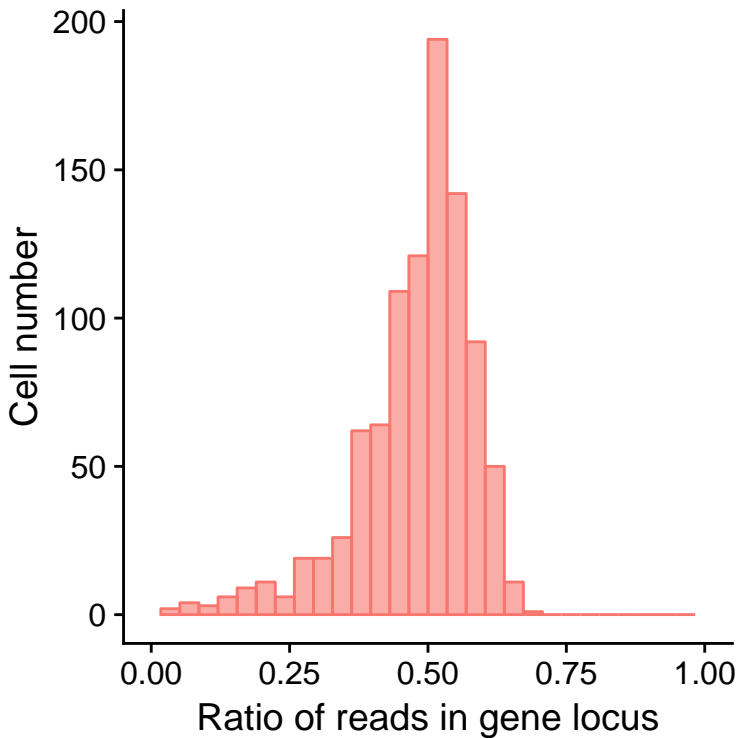
Uniquely mapped reads (Million)



Number of detected genes







Uniquely mapped reads (Million)

0

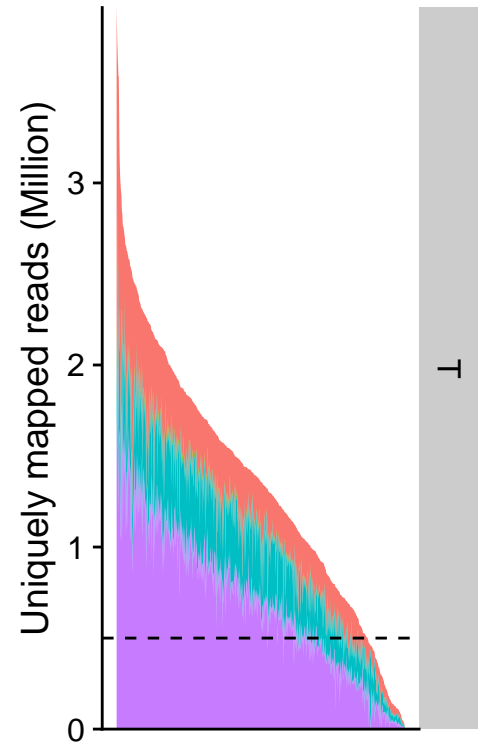
1

2

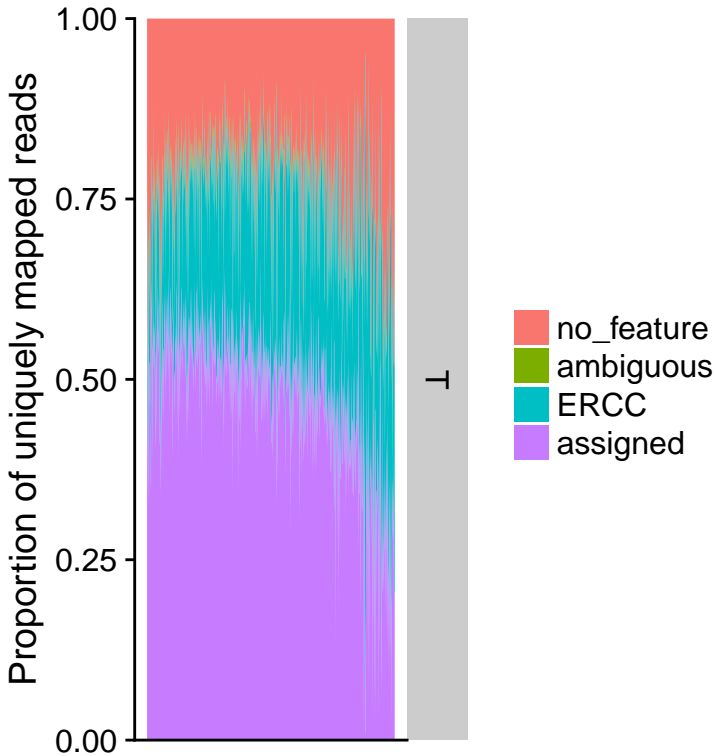
3

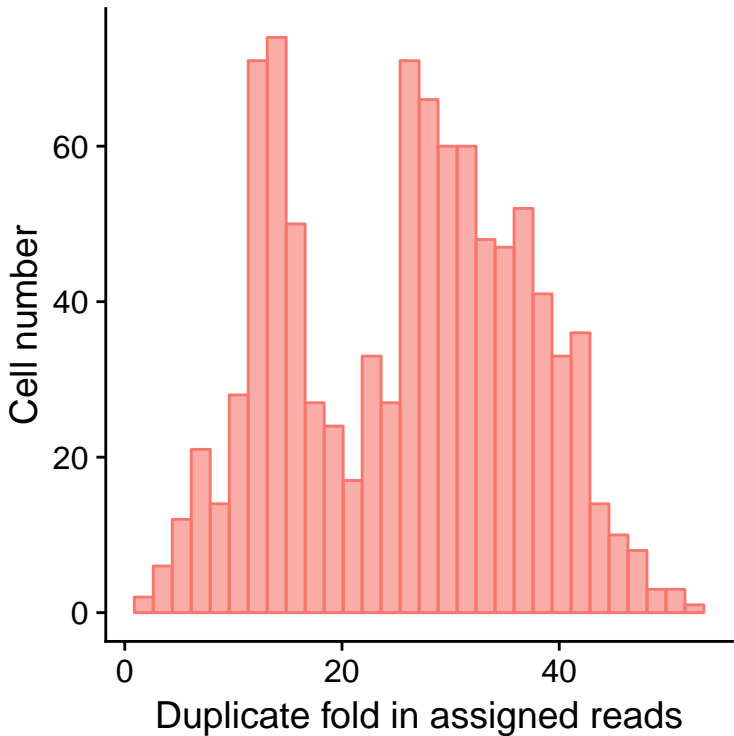
- no\_feature
- ambiguous
- ERCC
- assigned

1









Duplicate fold in ERCC reads

60  
40  
20  
0

0

10

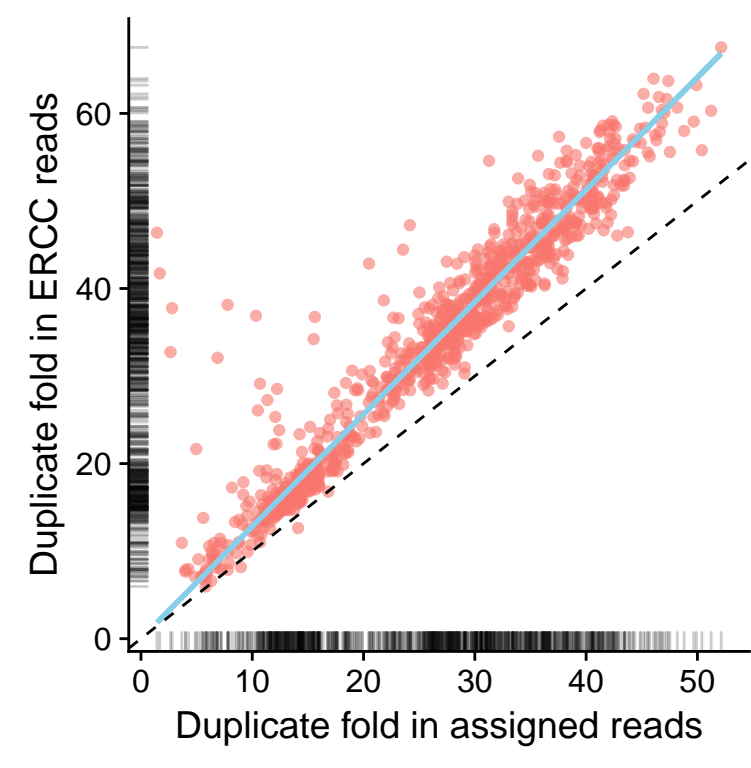
20

30

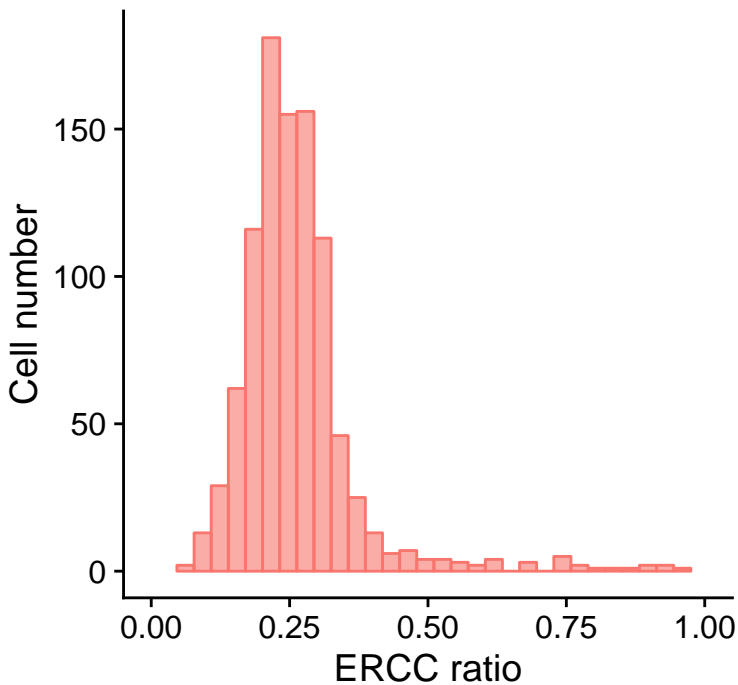
40

50

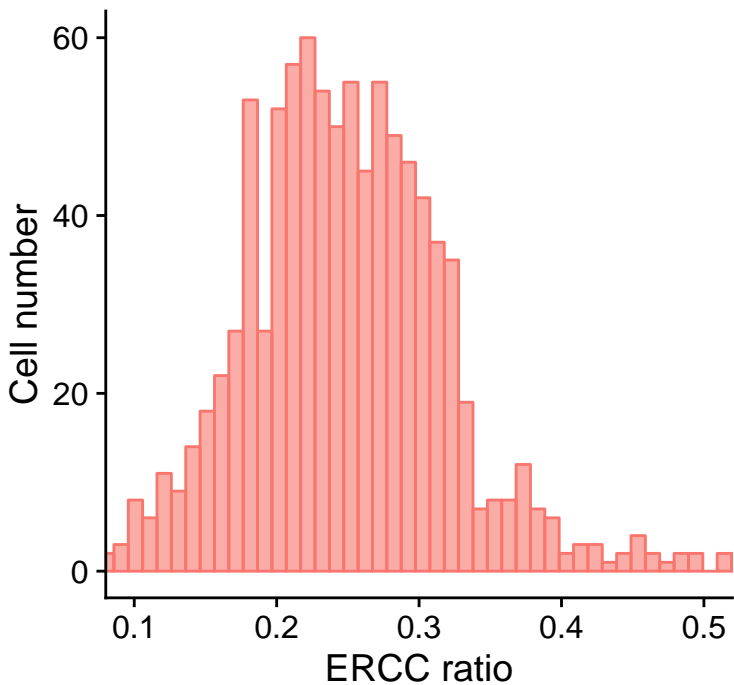
Duplicate fold in assigned reads



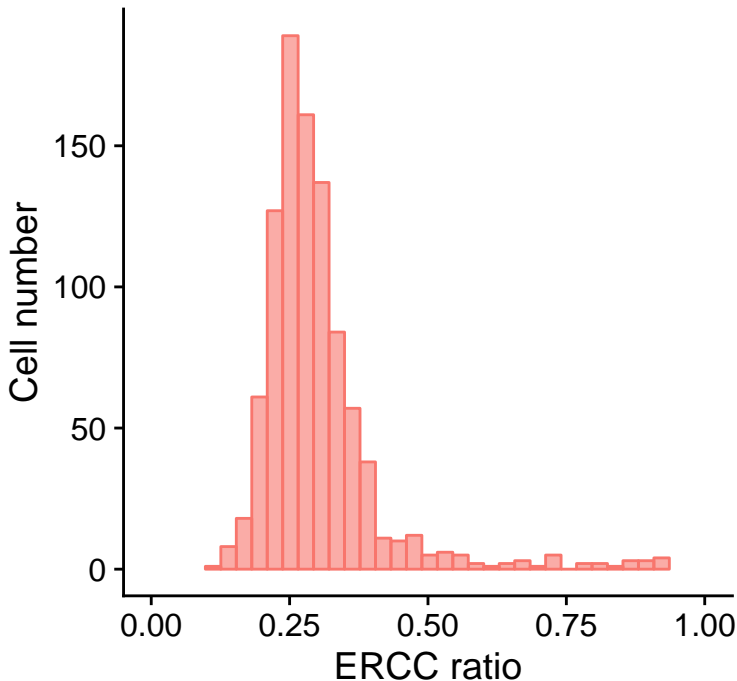
## Before removing duplicates



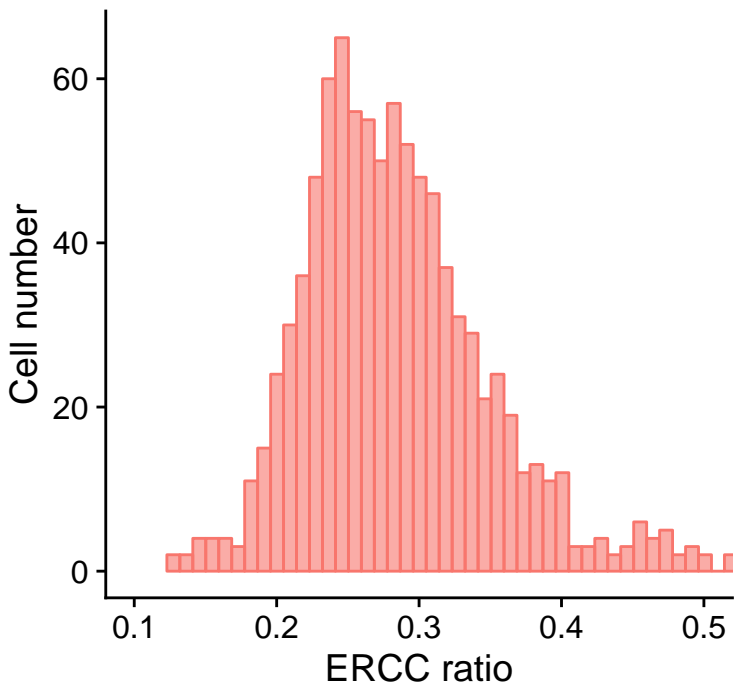
## Before removing dup. (Zoom in)



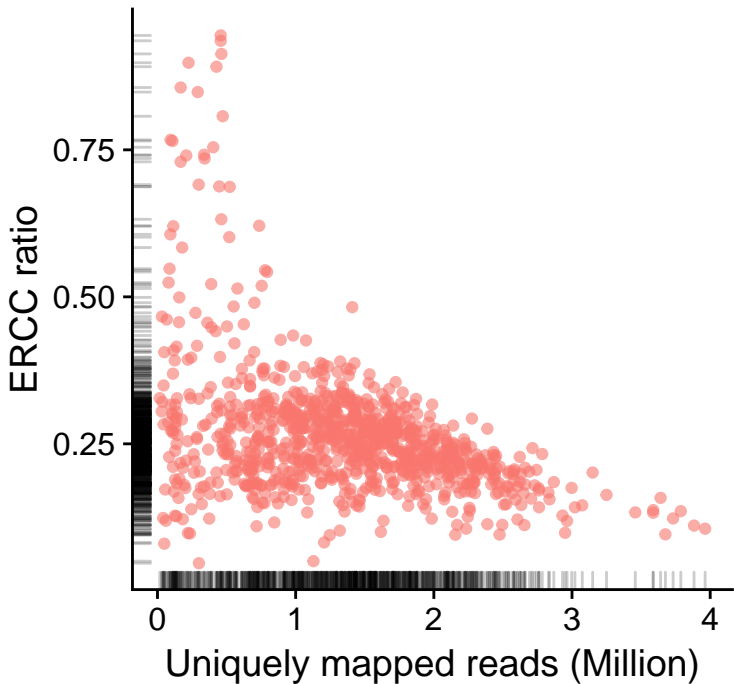
## After removing duplicates



## After removing dup. (Zoom in)



## Before removing duplicates





## After removing duplicates

