

Number of detected genes

6000

4000

2000

0

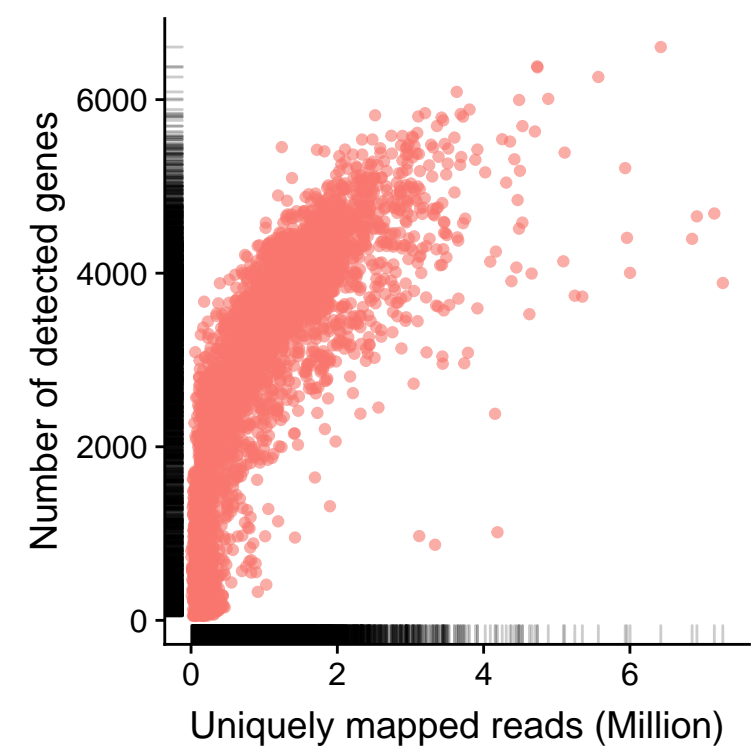
0

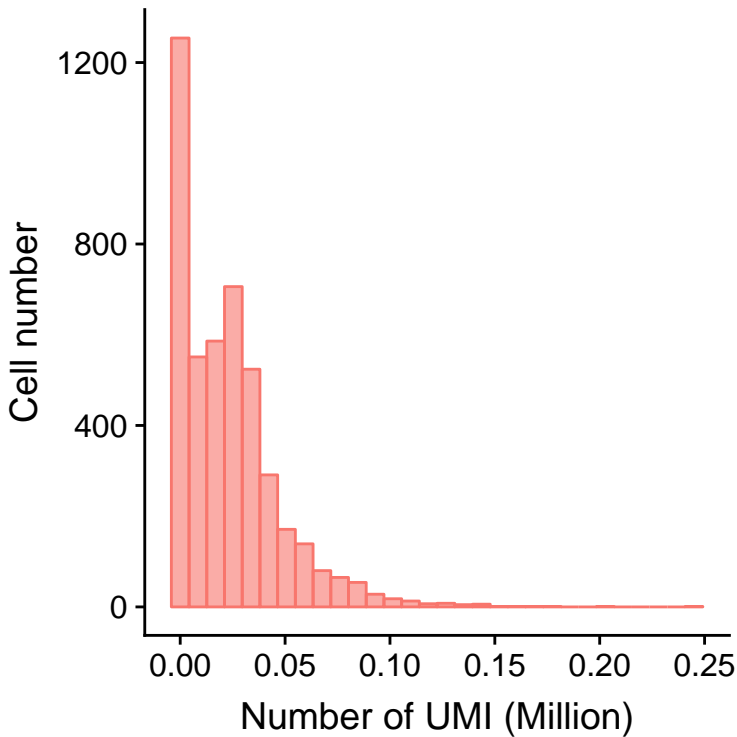
2

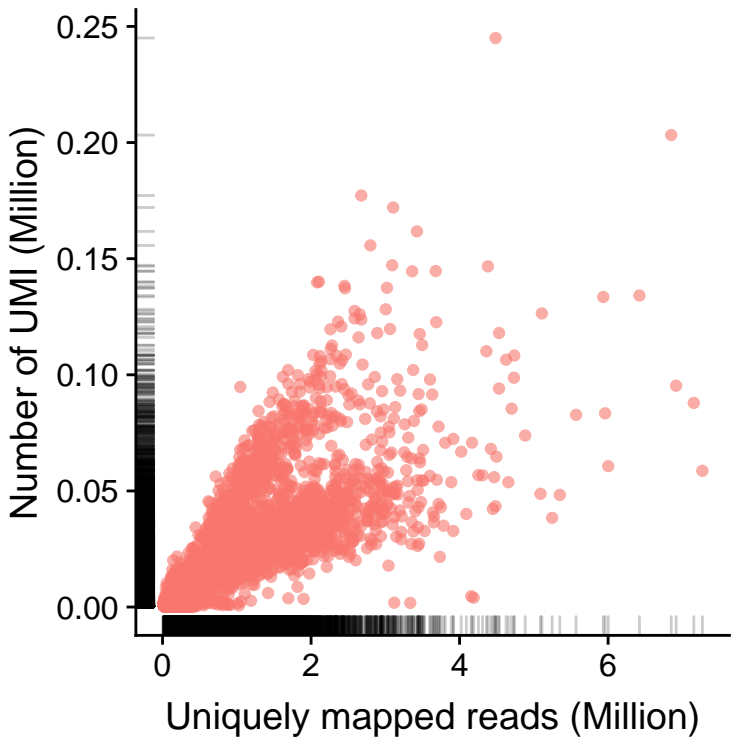
4

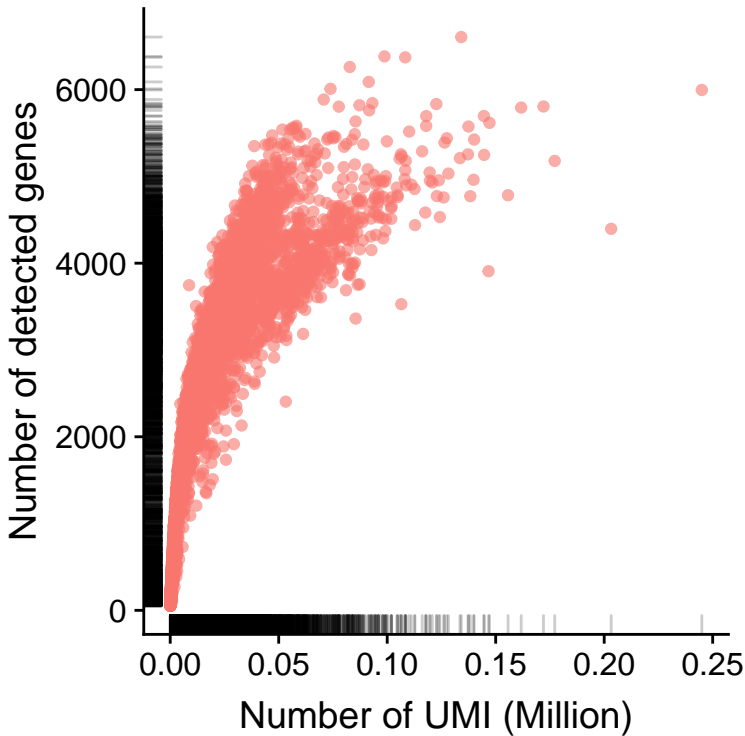
6

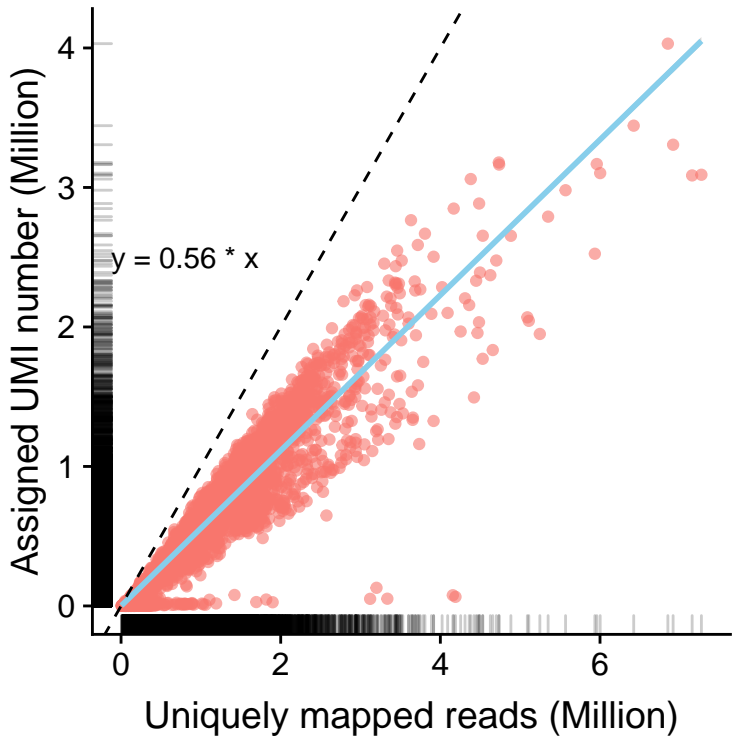
Uniquely mapped reads (Million)

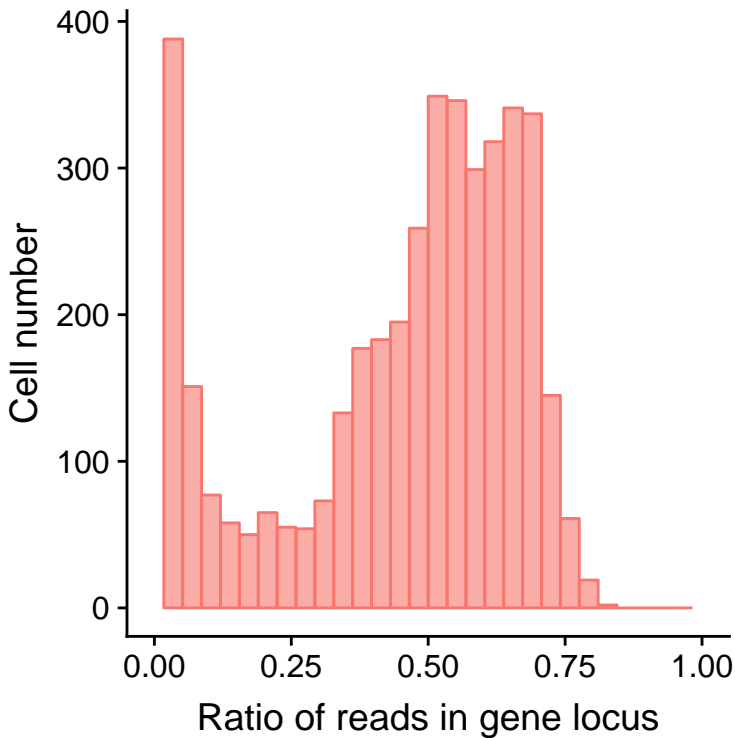




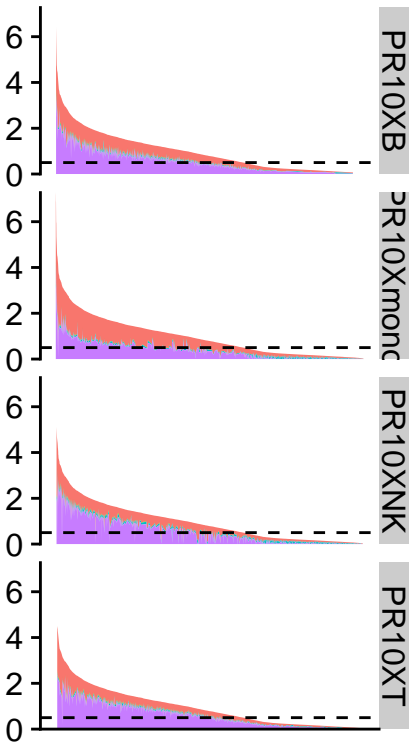






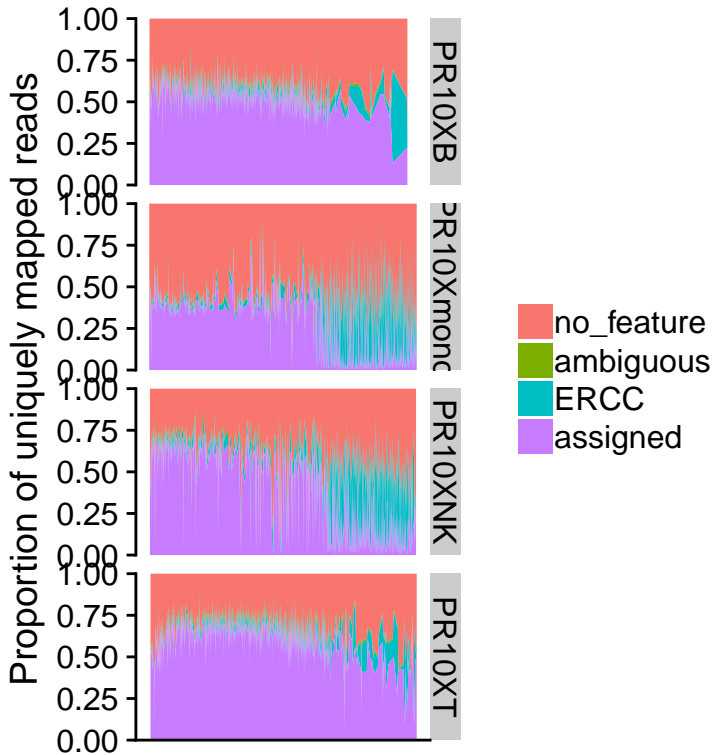


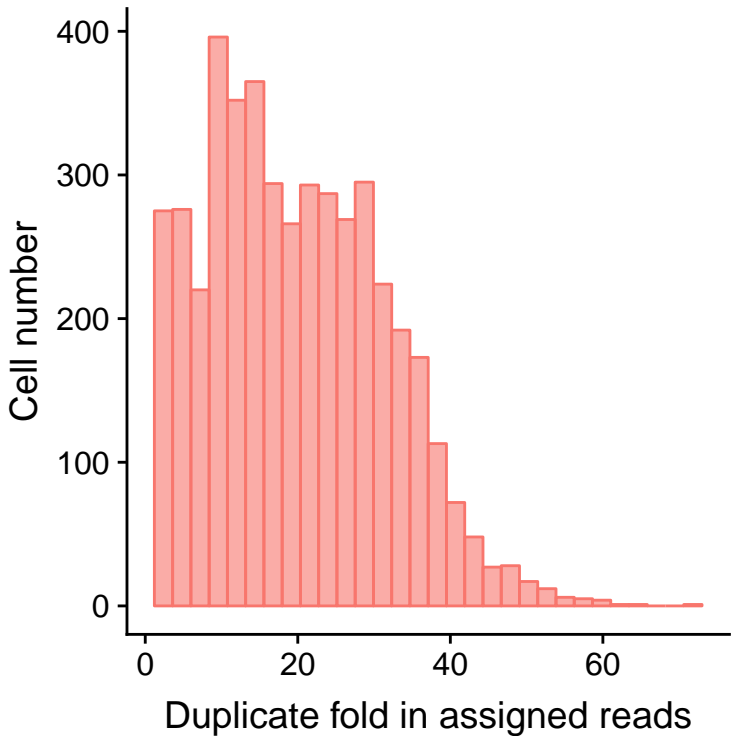
Uniquely mapped reads (Million)



no\_feature  
ambiguous  
ERCC  
assigned







Duplicate fold in ERCC reads

75

50

25

0

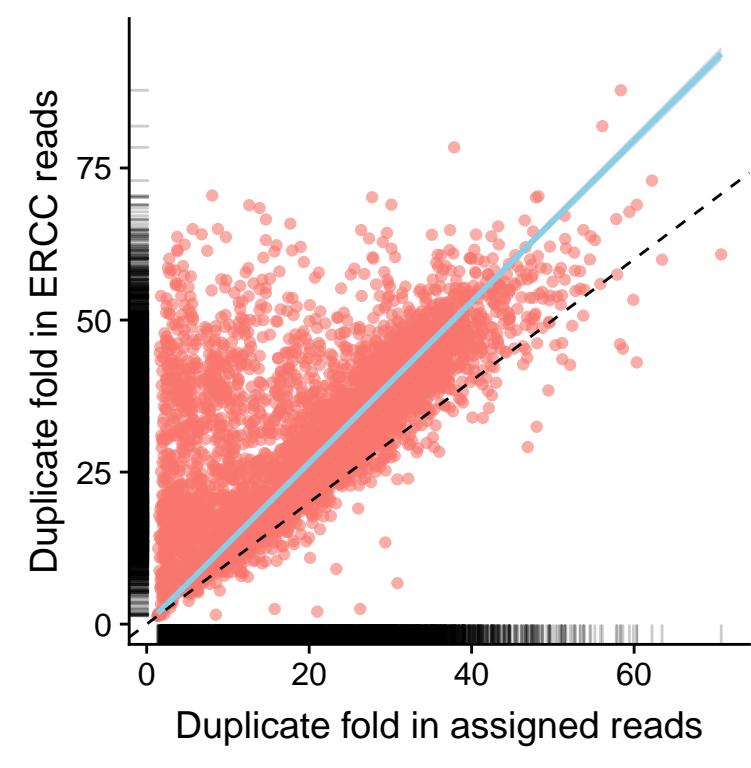
0

20

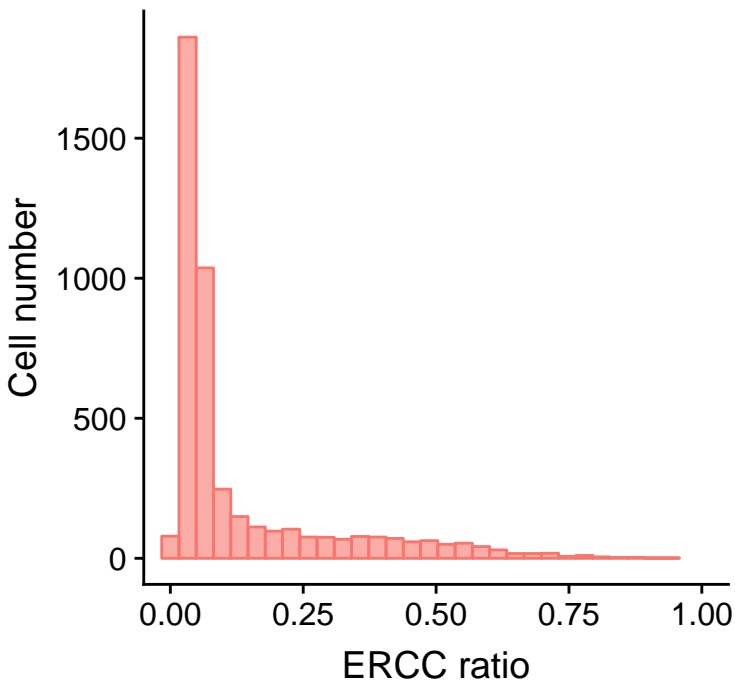
40

60

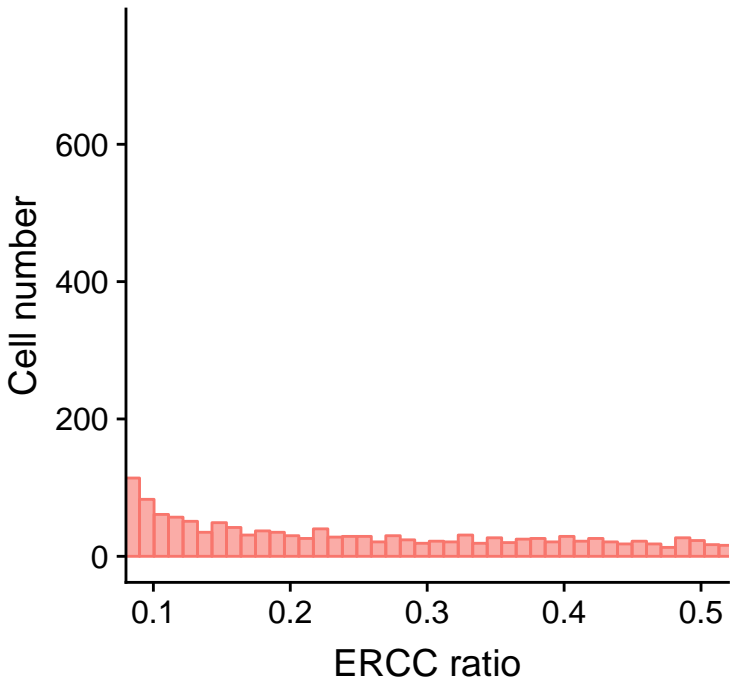
Duplicate fold in assigned reads



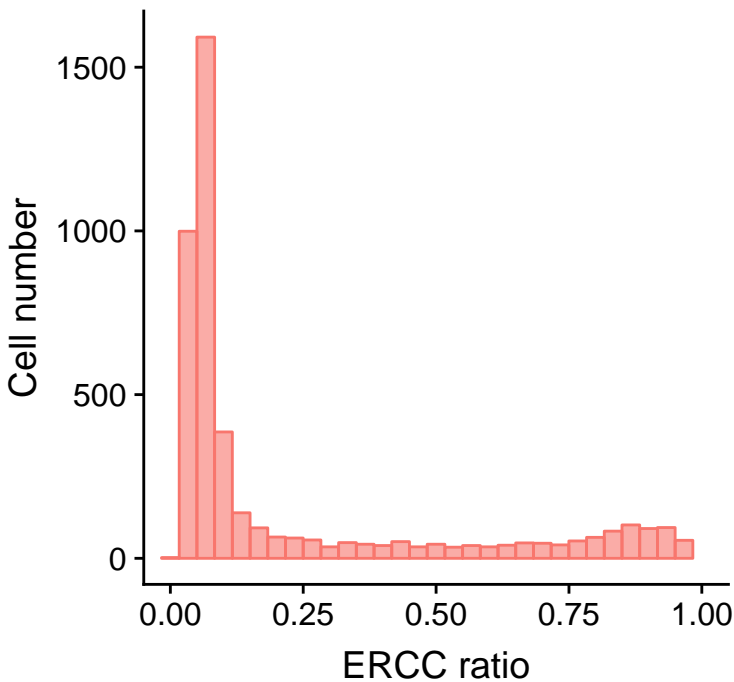
## Before removing duplicates



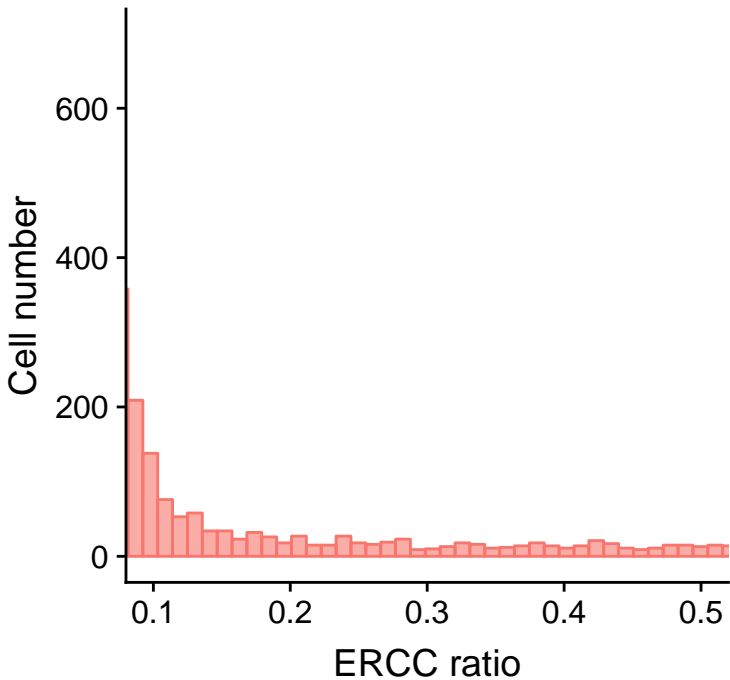
## Before removing dup. (Zoom in)



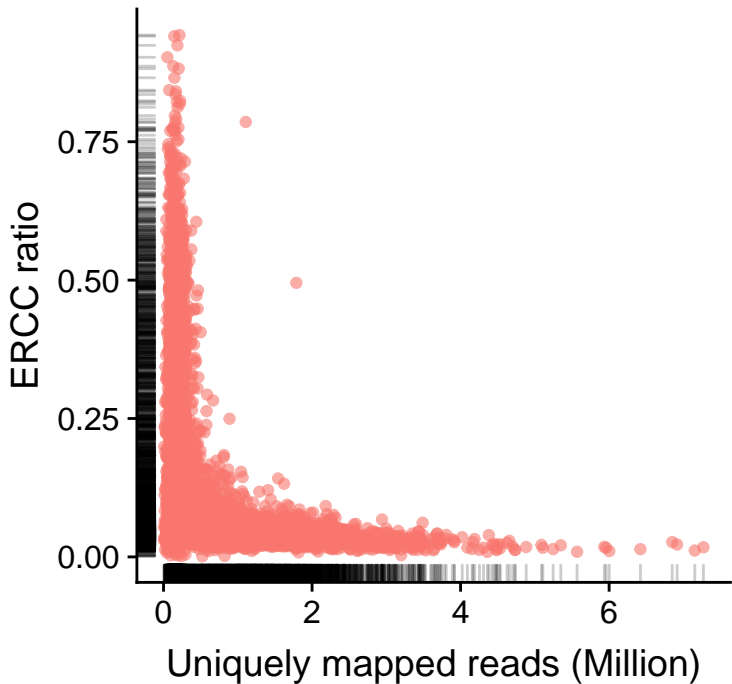
## After removing duplicates



## After removing dup. (Zoom in)



## Before removing duplicates





## After removing duplicates

