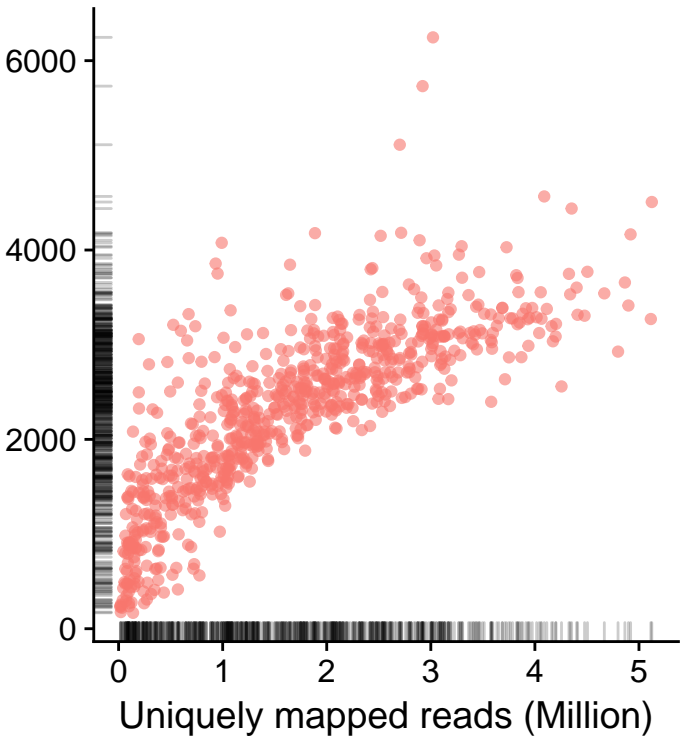
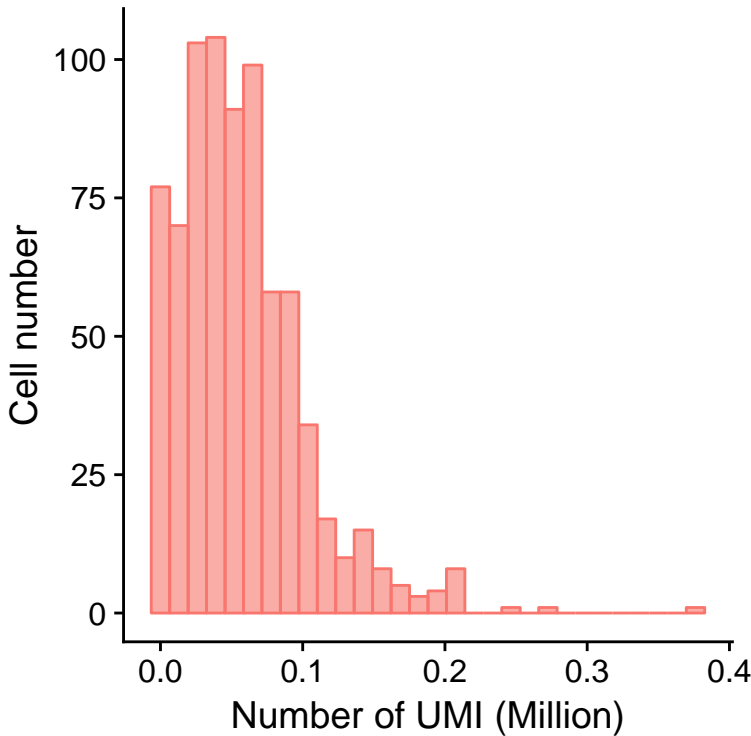


Number of detected genes





Number of UMI (Million)

0.3
0.2
0.1
0.0

0

1

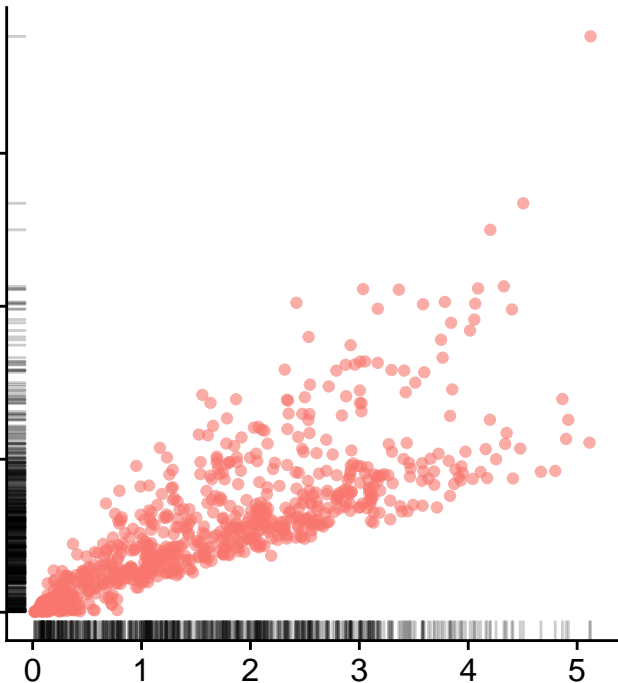
2

3

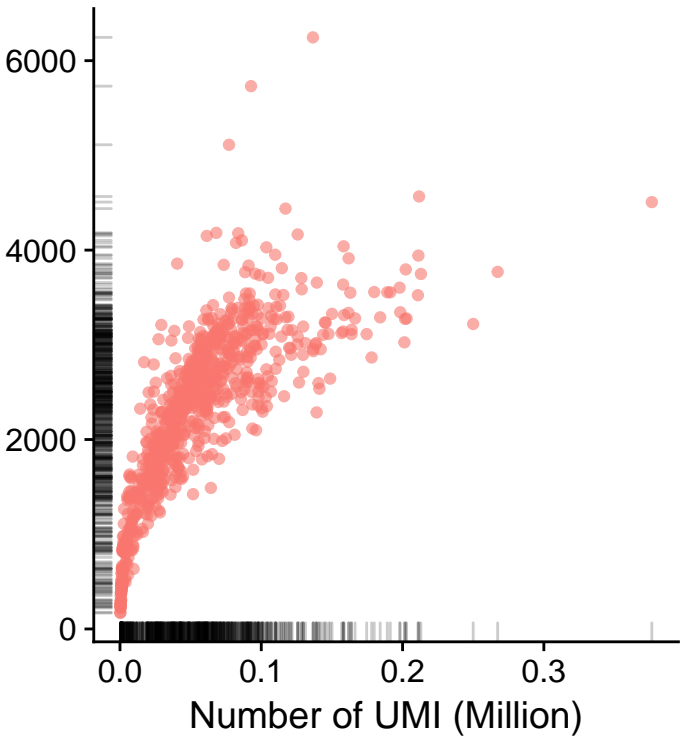
4

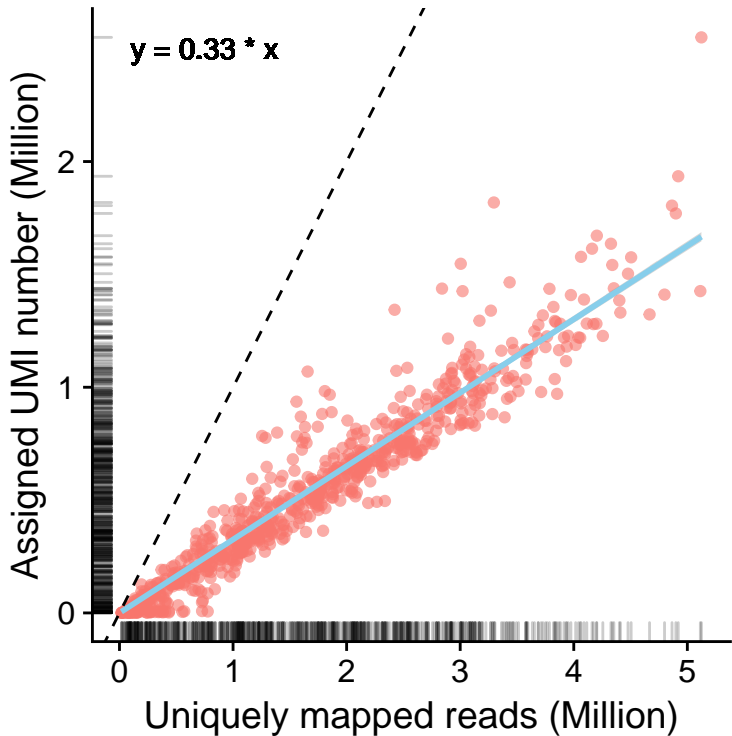
5

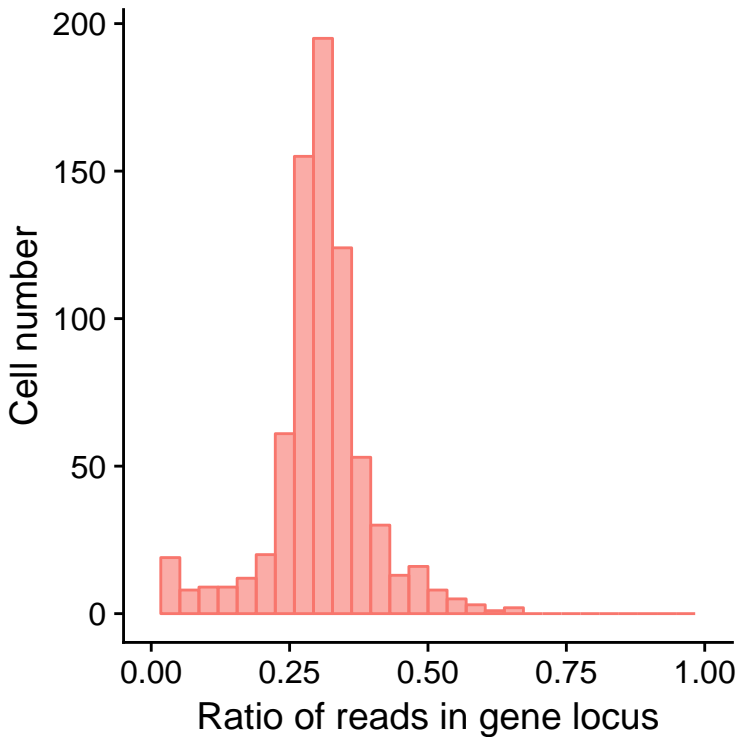
Uniquely mapped reads (Million)

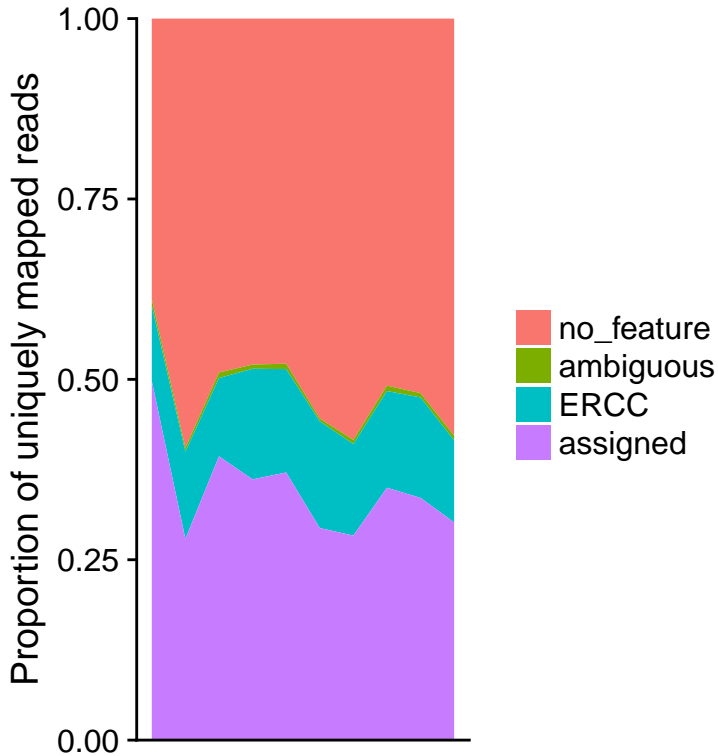


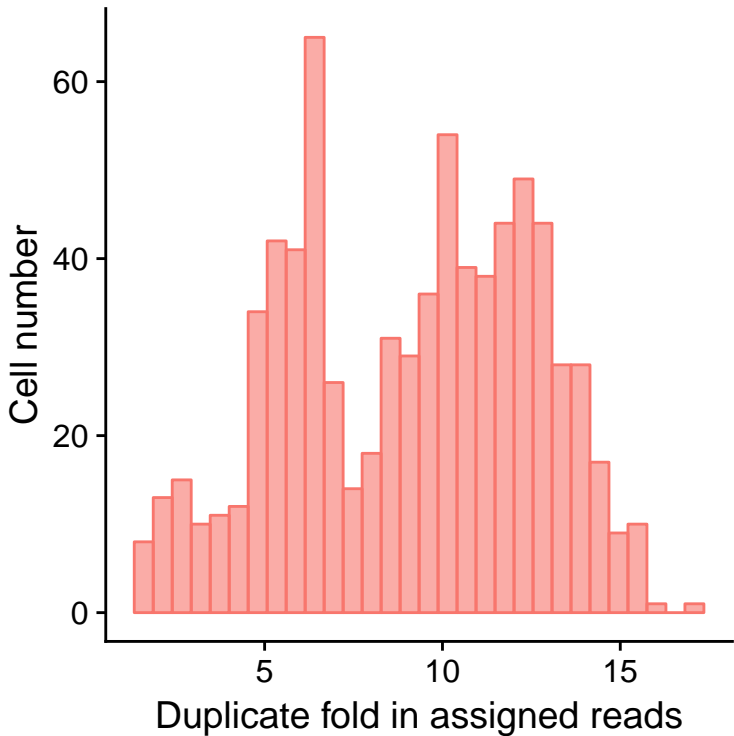
Number of detected genes











Duplicate fold in ERCC reads

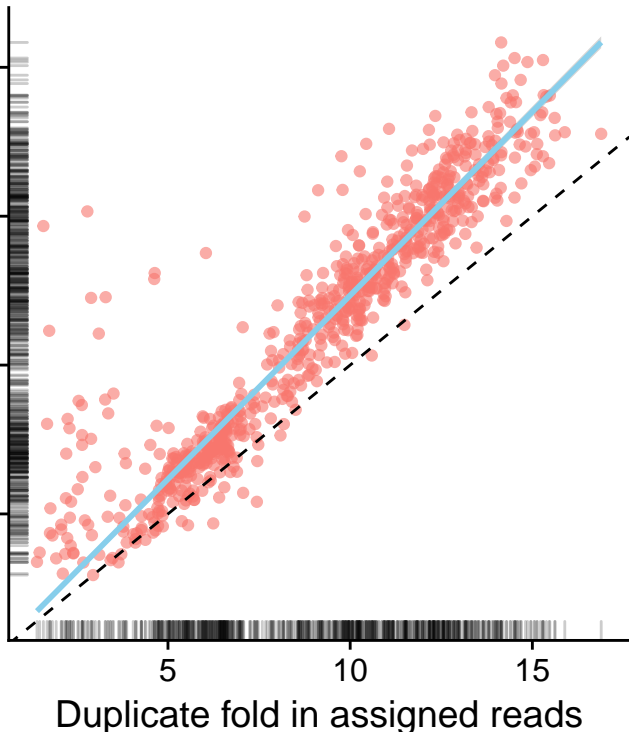
20
15
10
5

5

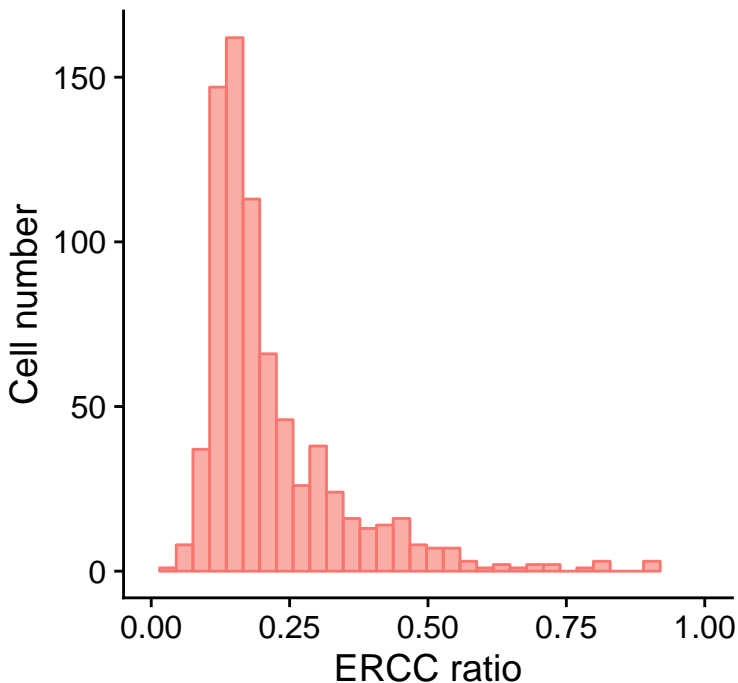
10

15

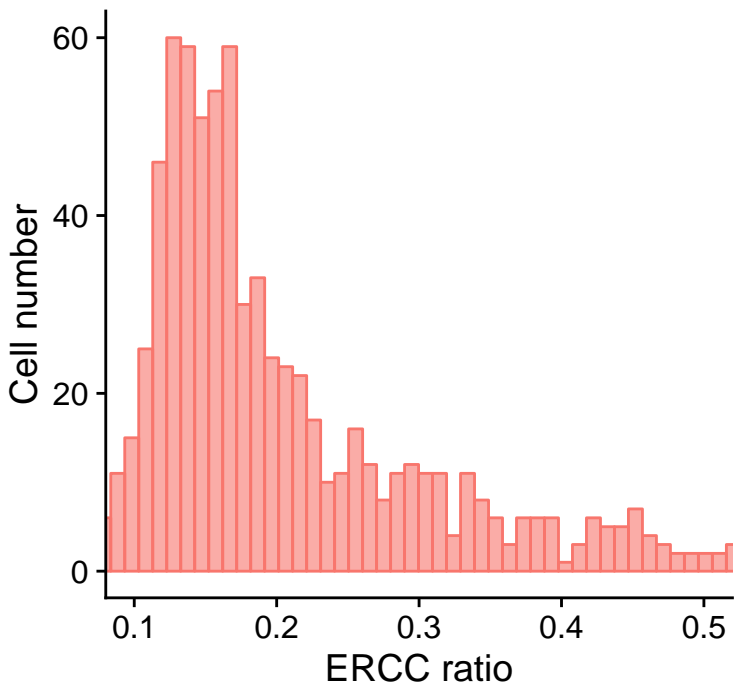
Duplicate fold in assigned reads



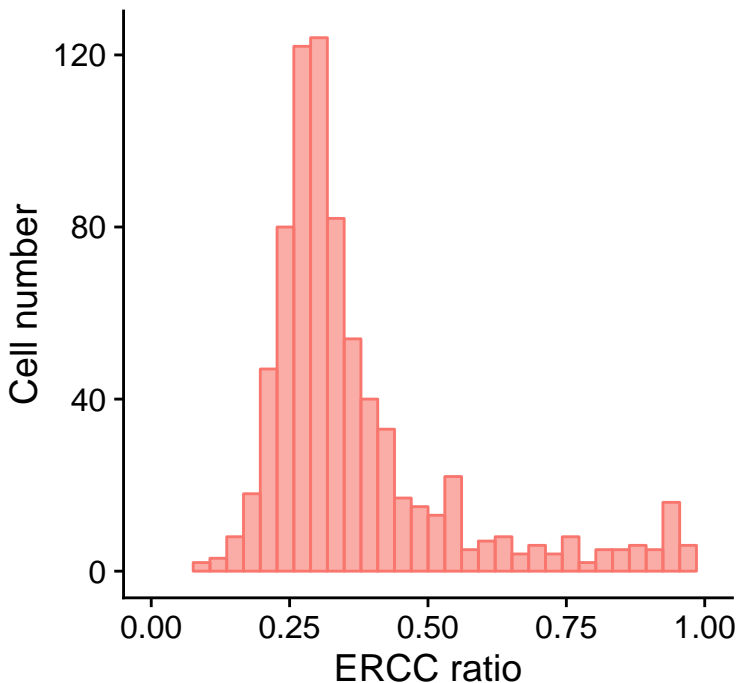
Before removing duplicates



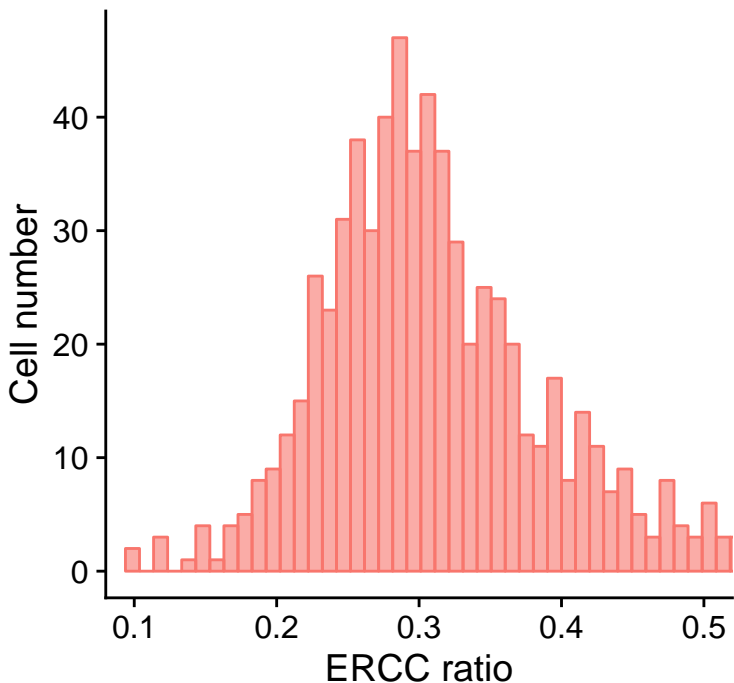
Before removing dup. (Zoom in)



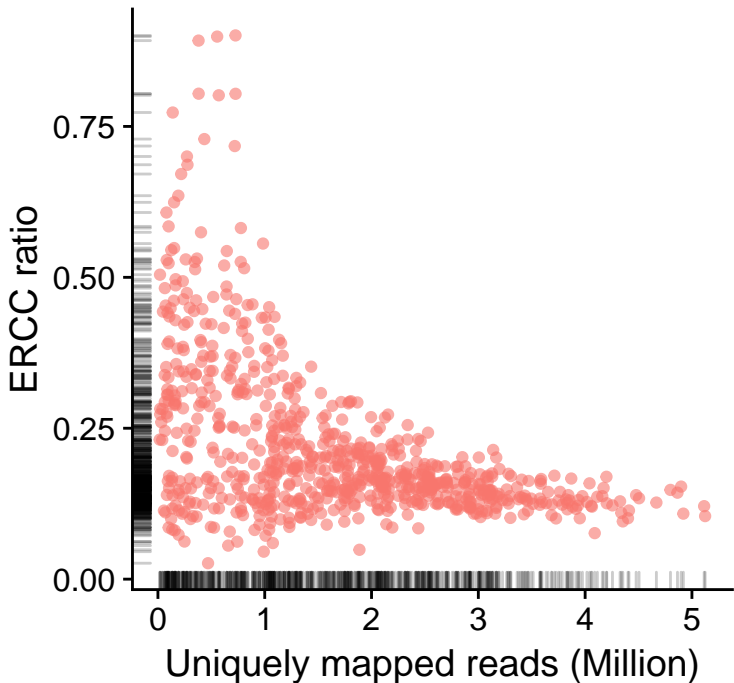
After removing duplicates



After removing dup. (Zoom in)



Before removing duplicates



After removing duplicates

