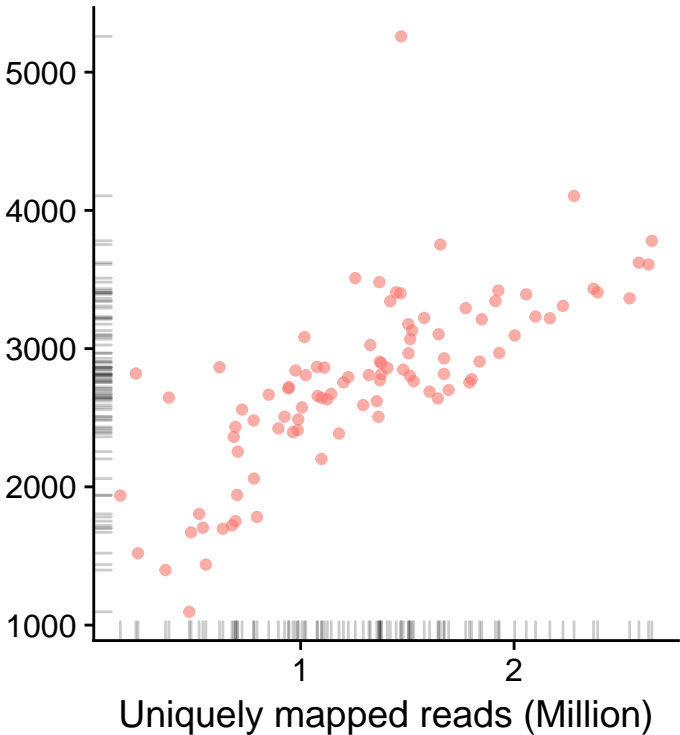
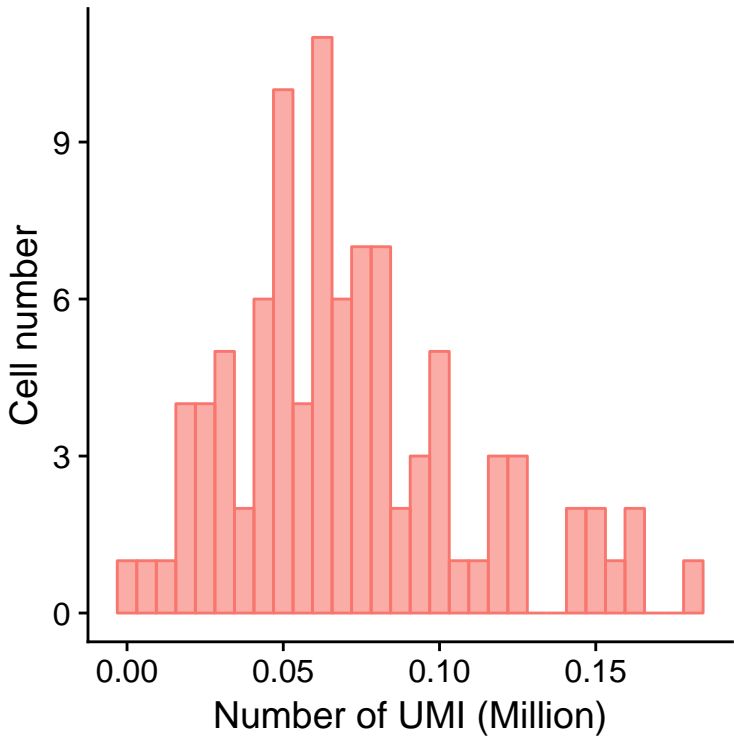


Number of detected genes





Number of UMI (Million)

0.15

0.10

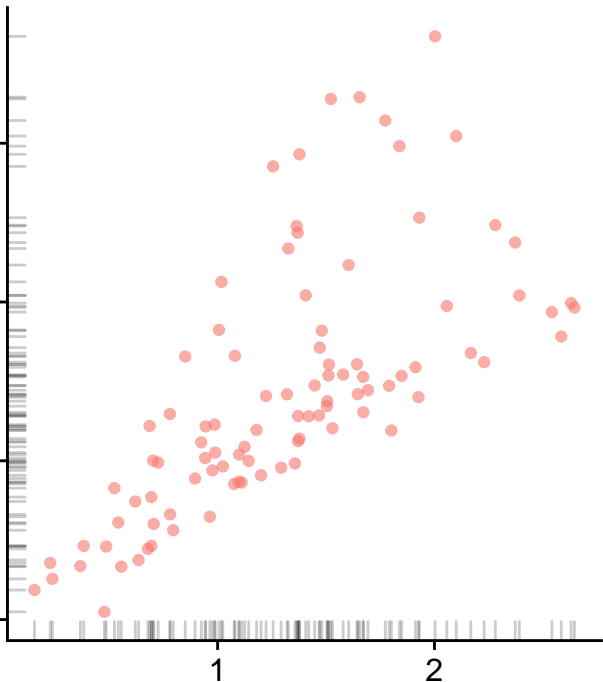
0.05

0.00

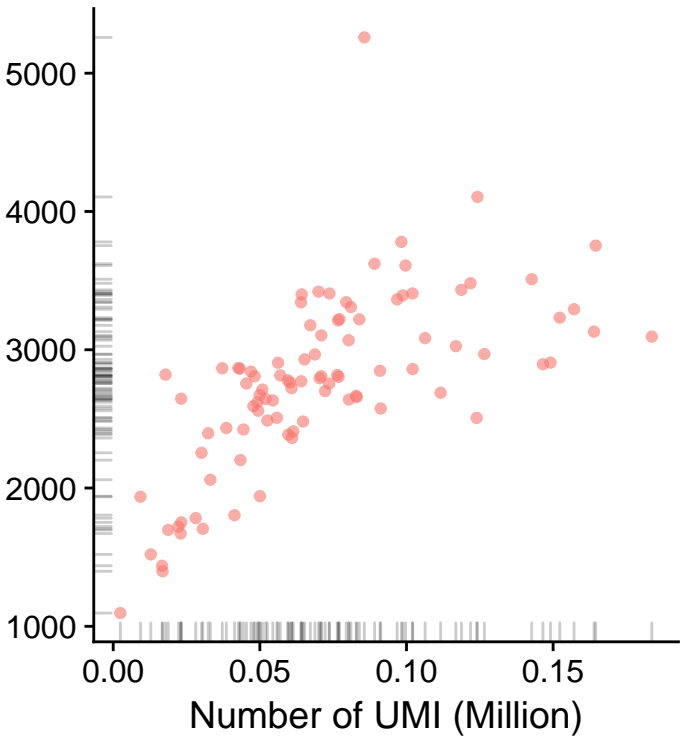
1

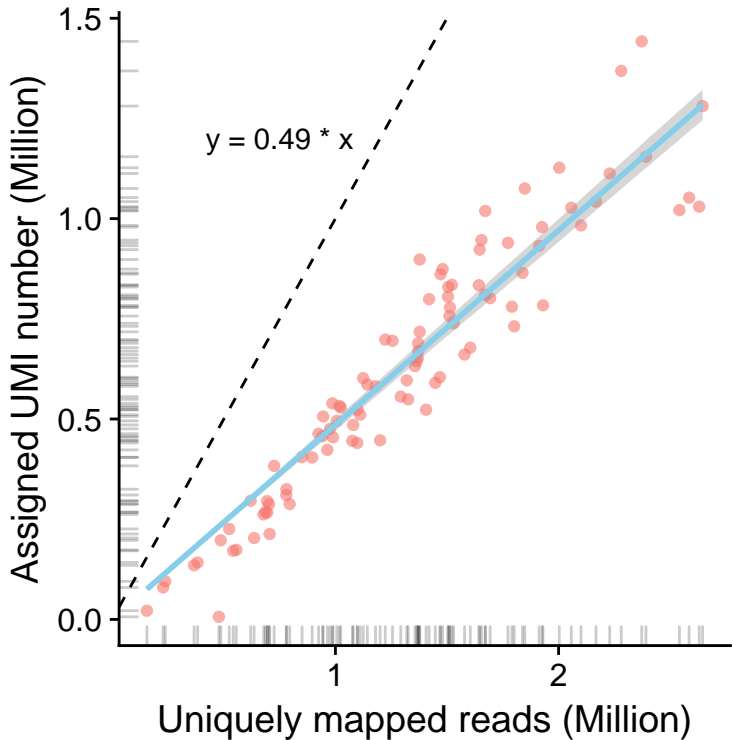
2

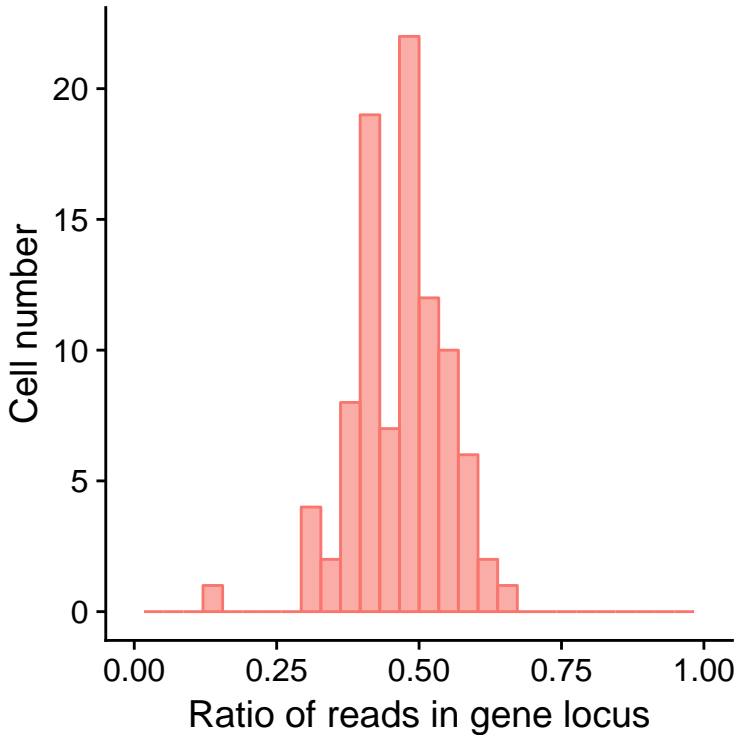
Uniquely mapped reads (Million)

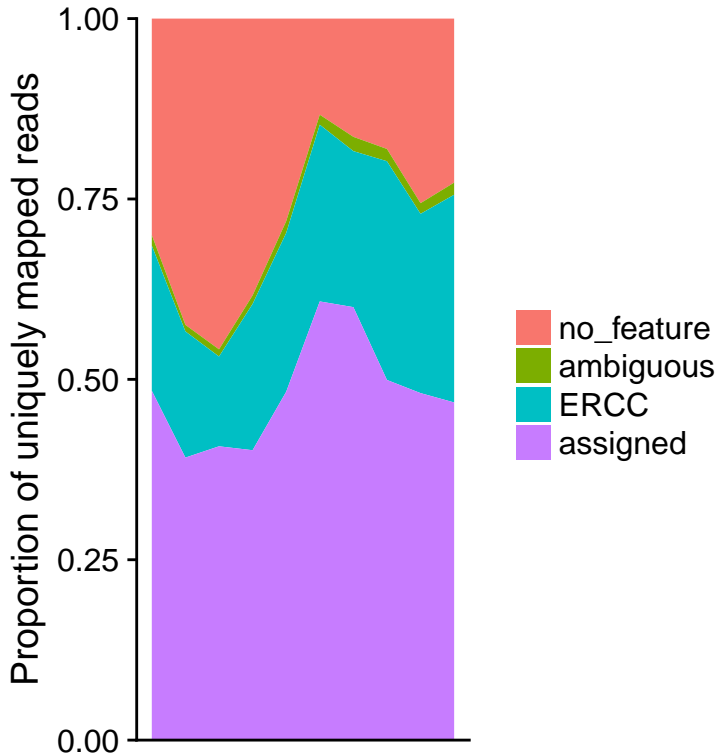


Number of detected genes

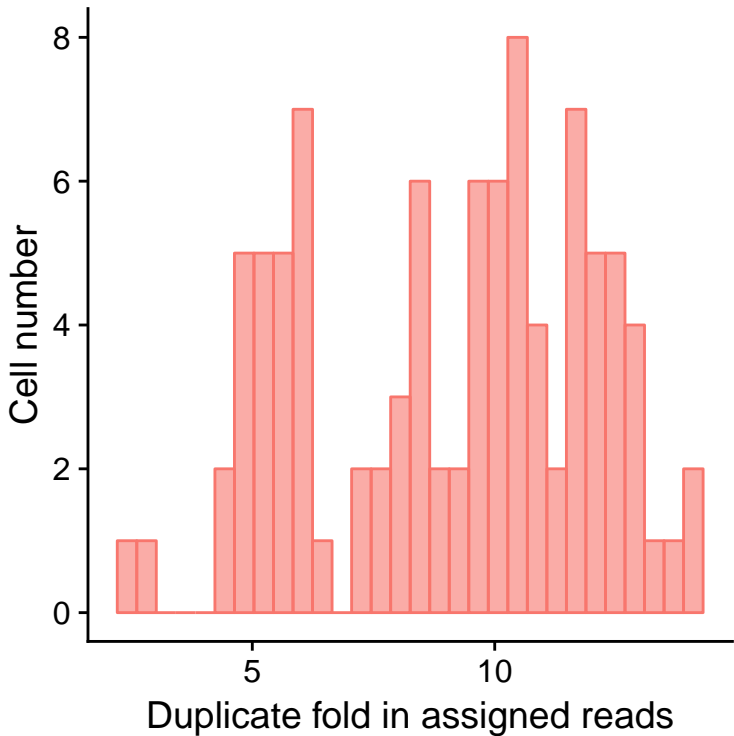












Duplicate fold in ERCC reads

20

15

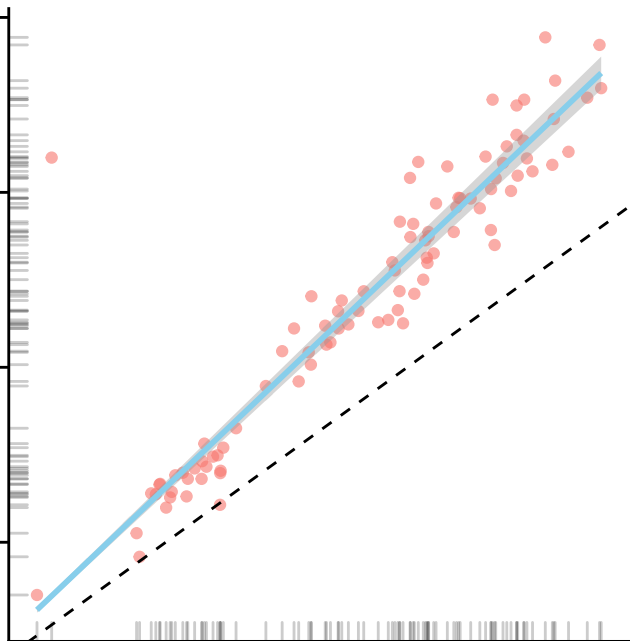
10

5

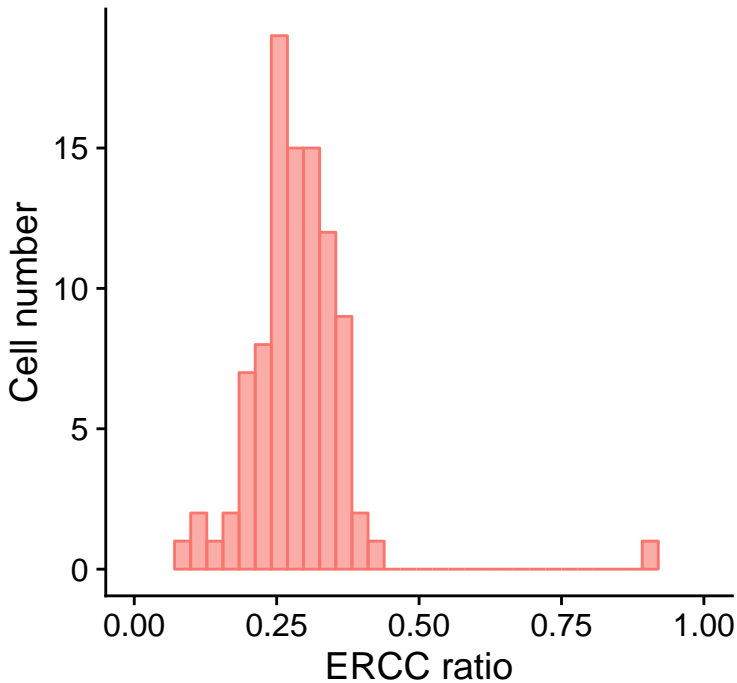
5

10

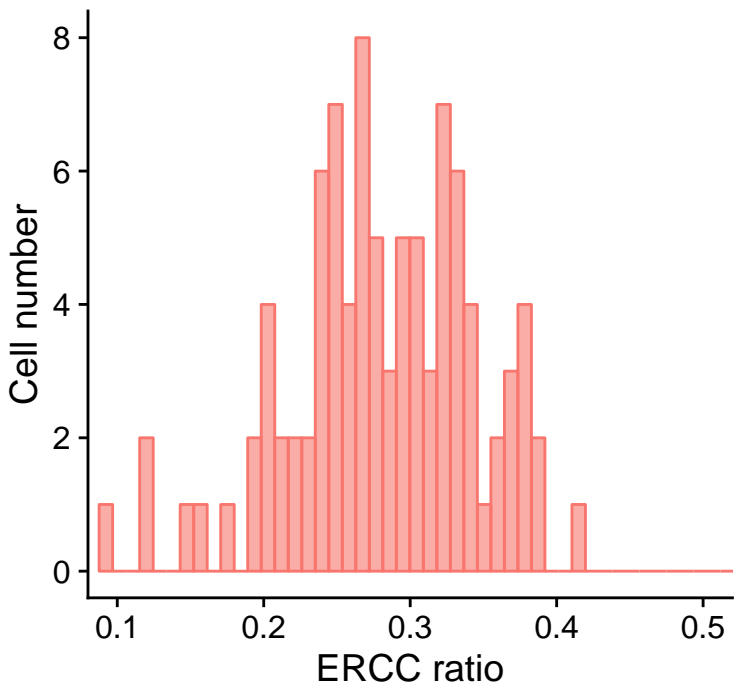
Duplicate fold in assigned reads



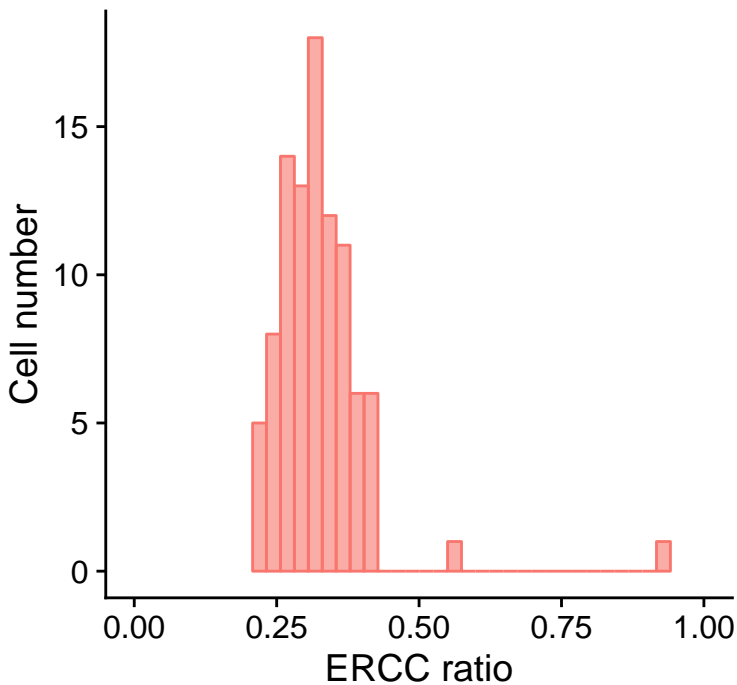
## Before removing duplicates



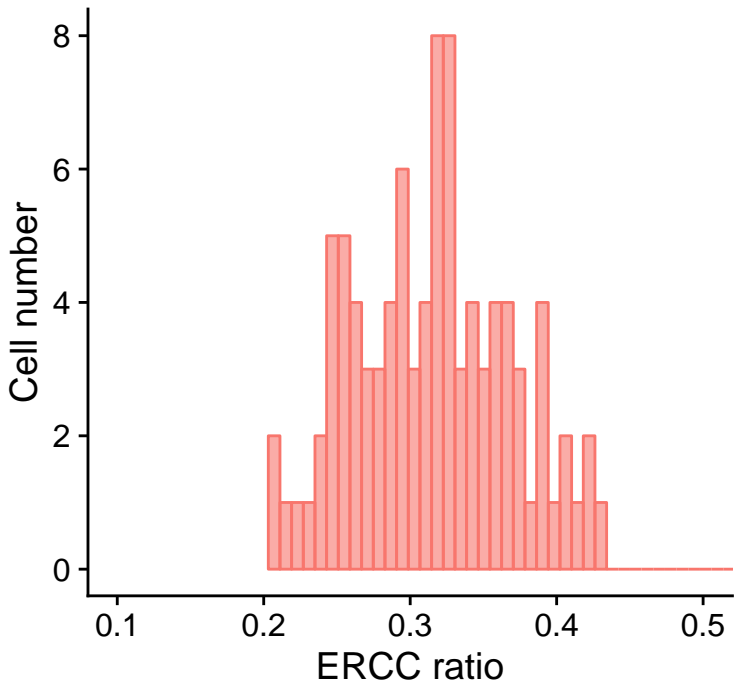
# Before removing dup. (Zoom in)



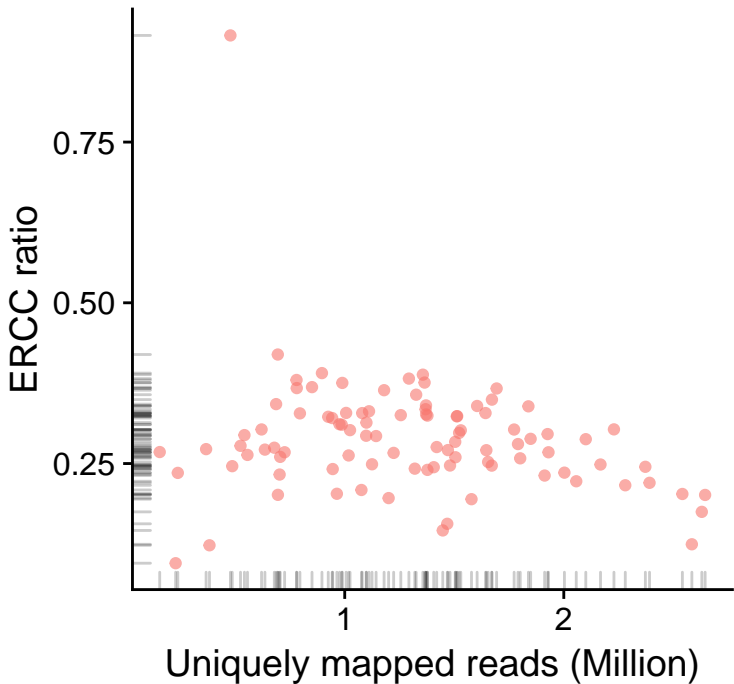
## After removing duplicates



## After removing dup. (Zoom in)



## Before removing duplicates



## After removing duplicates

