

Number of detected genes

10000

8000

6000

0.0

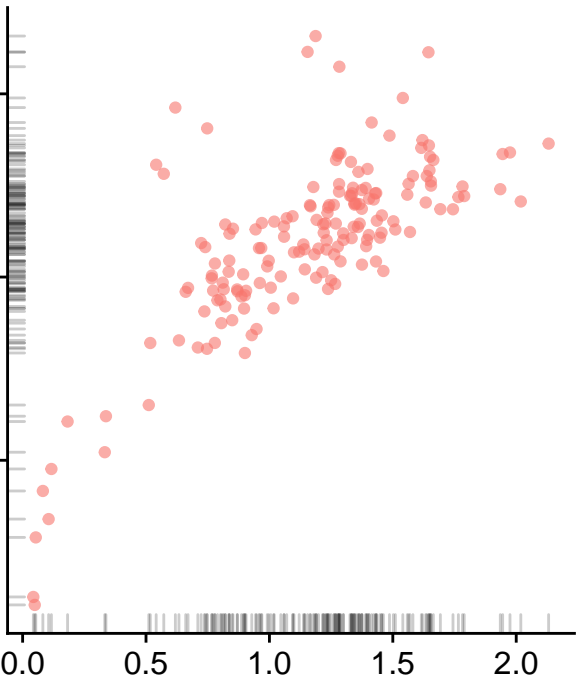
0.5

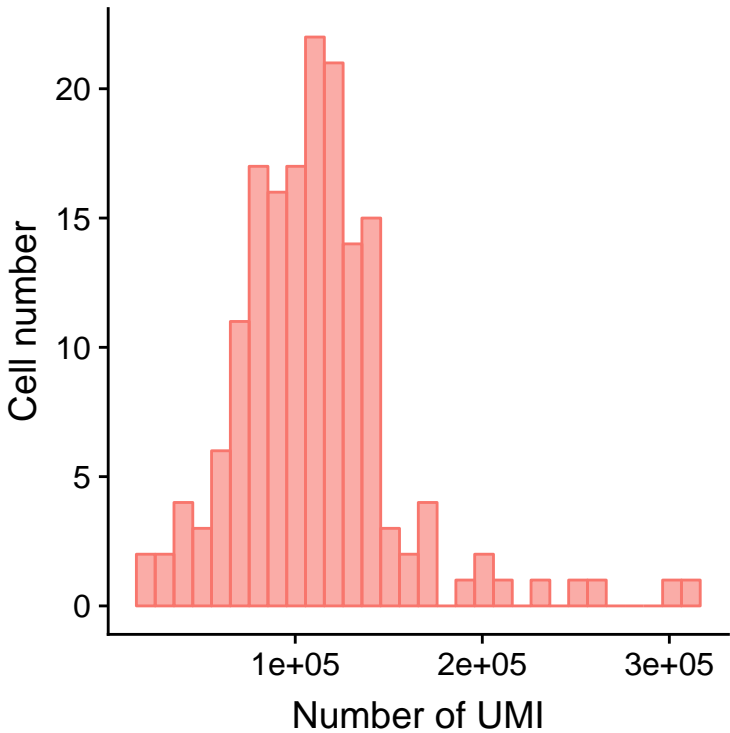
1.0

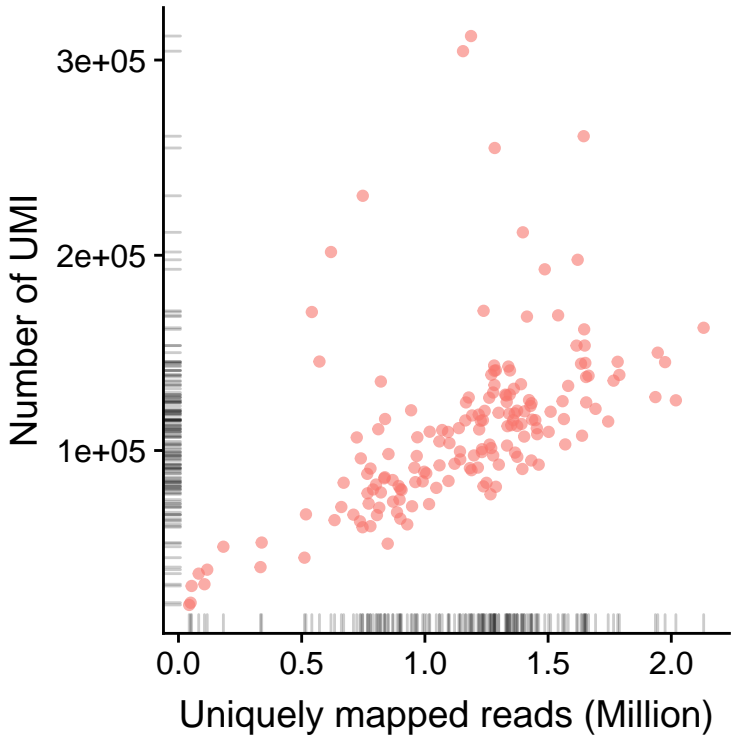
1.5

2.0

Uniquely mapped reads (Million)







Number of detected genes

10000

8000

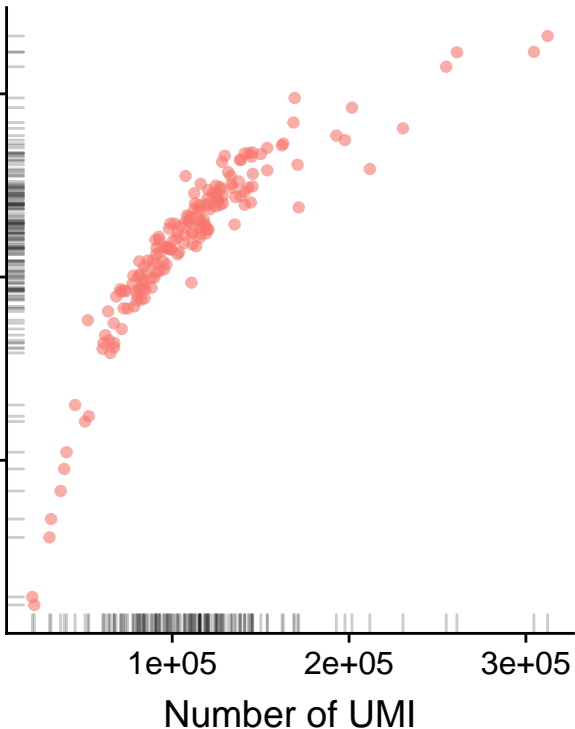
6000

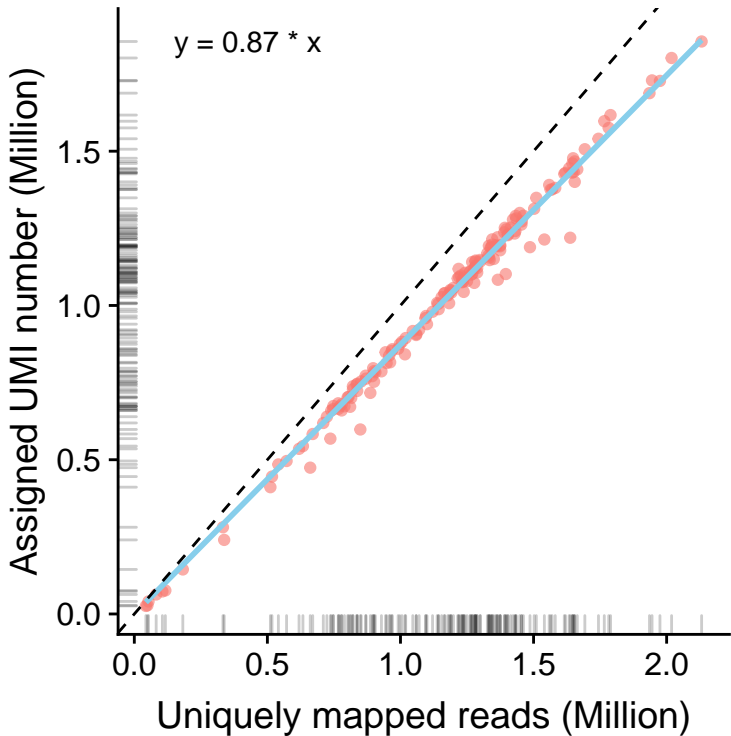
1e+05

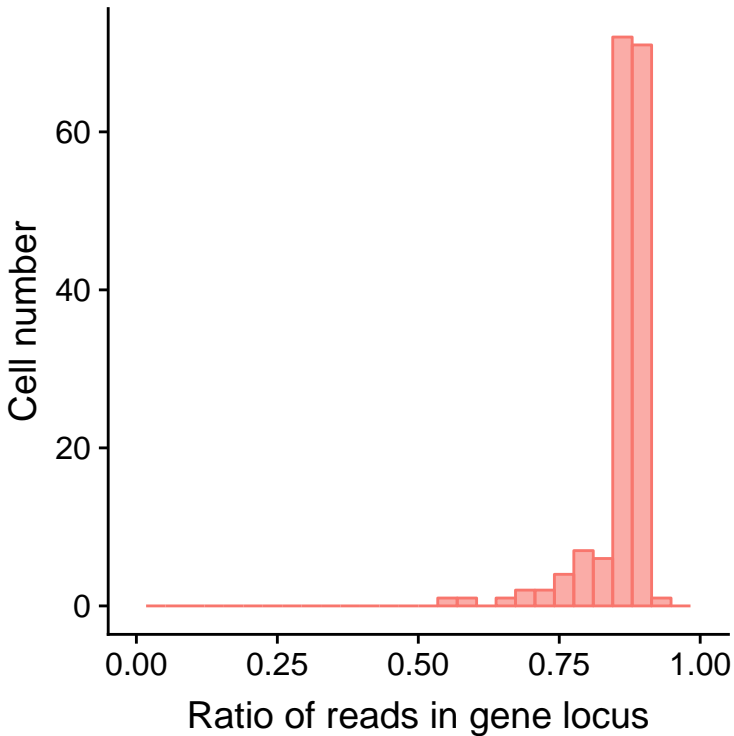
2e+05

3e+05

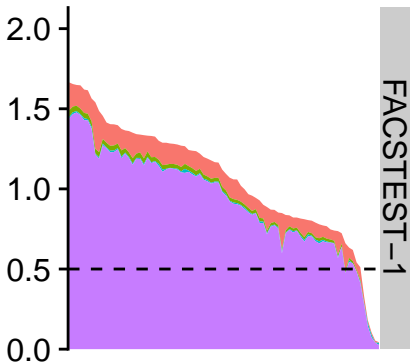
Number of UMI



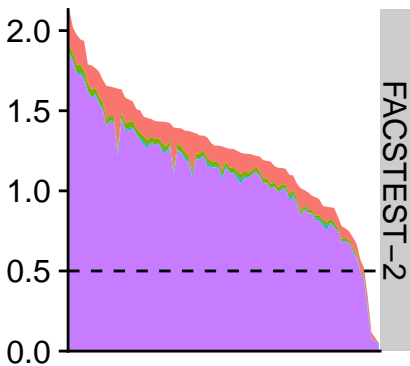


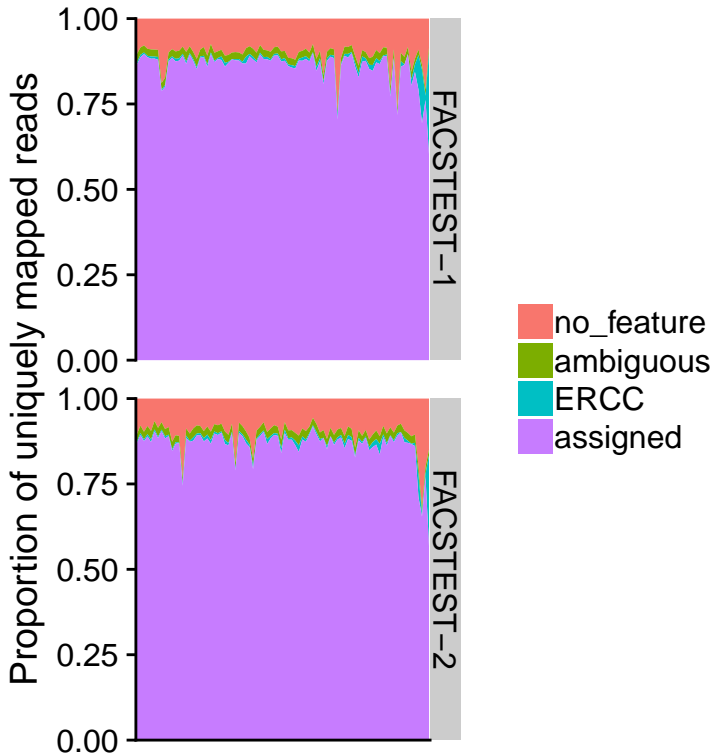


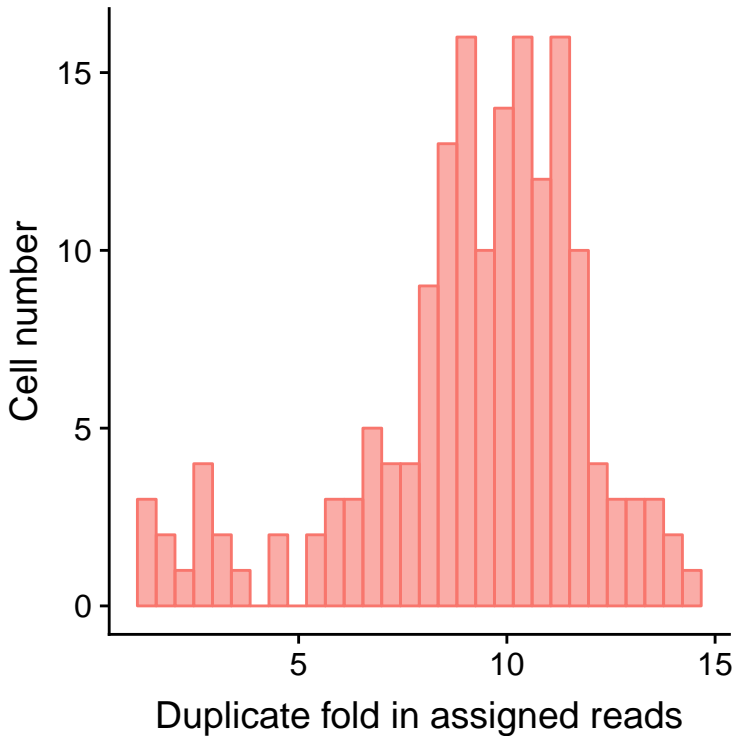
Uniquely mapped reads (Million)



no_feature
ambiguous
ERCC
assigned







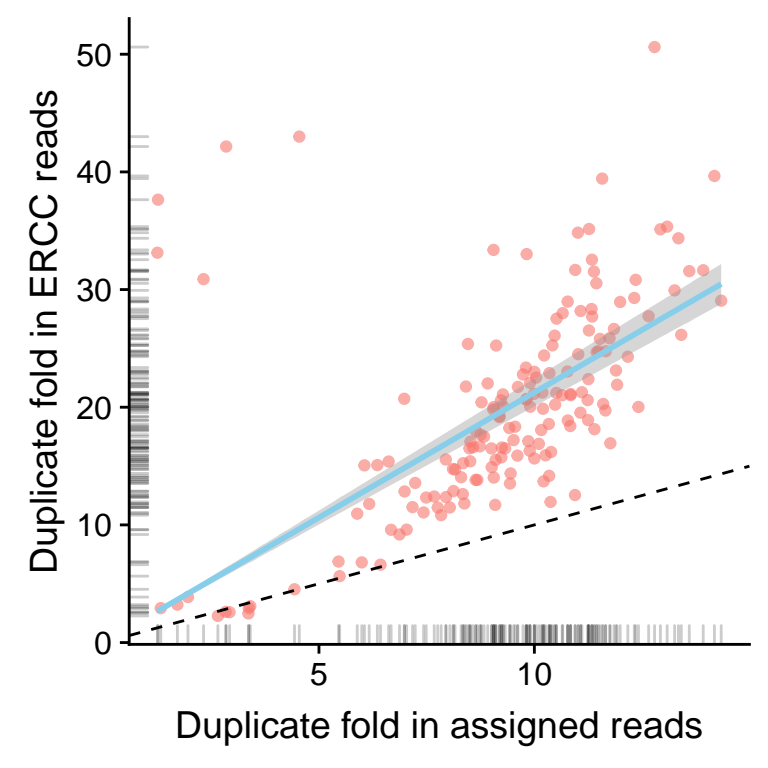
Duplicate fold in ERCC reads

50
40
30
20
10
0

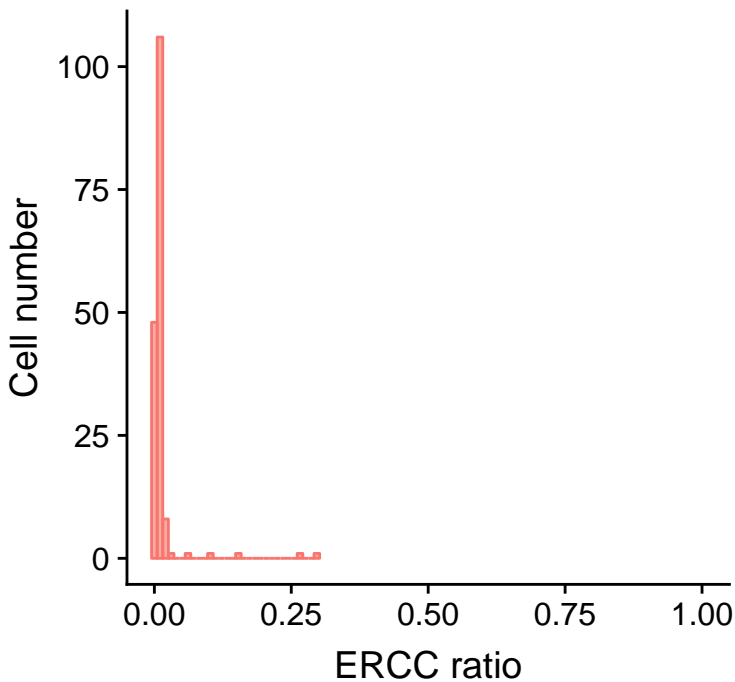
5

10

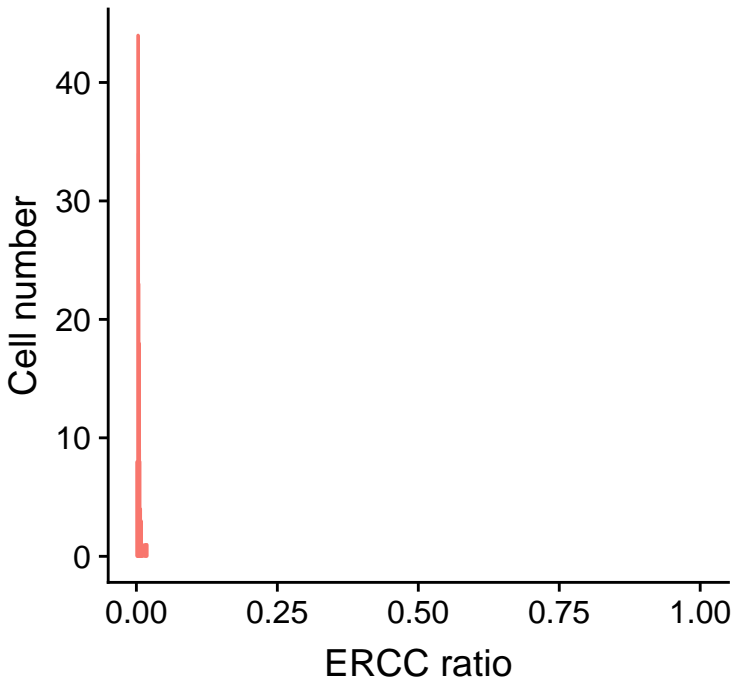
Duplicate fold in assigned reads



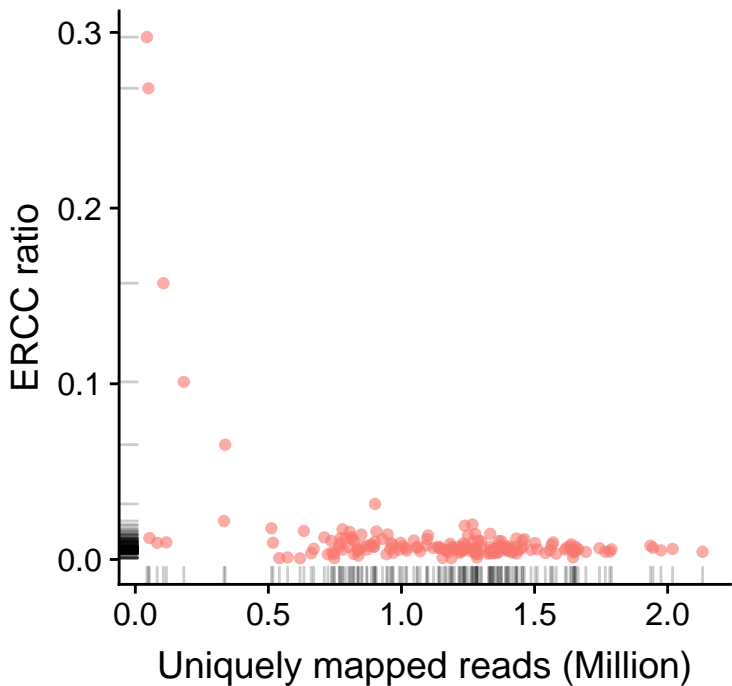
Before removing duplicates



After removing duplicates



Before removing duplicates



After removing duplicates

