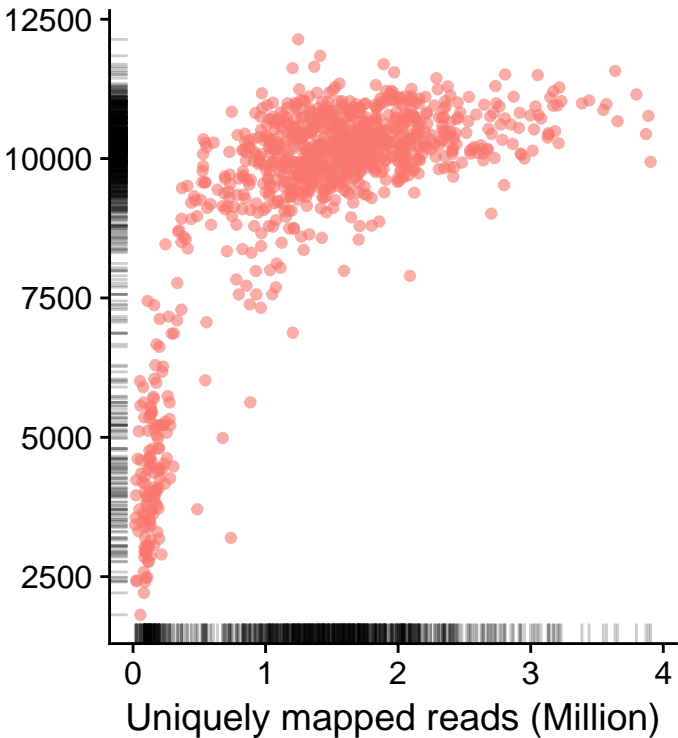
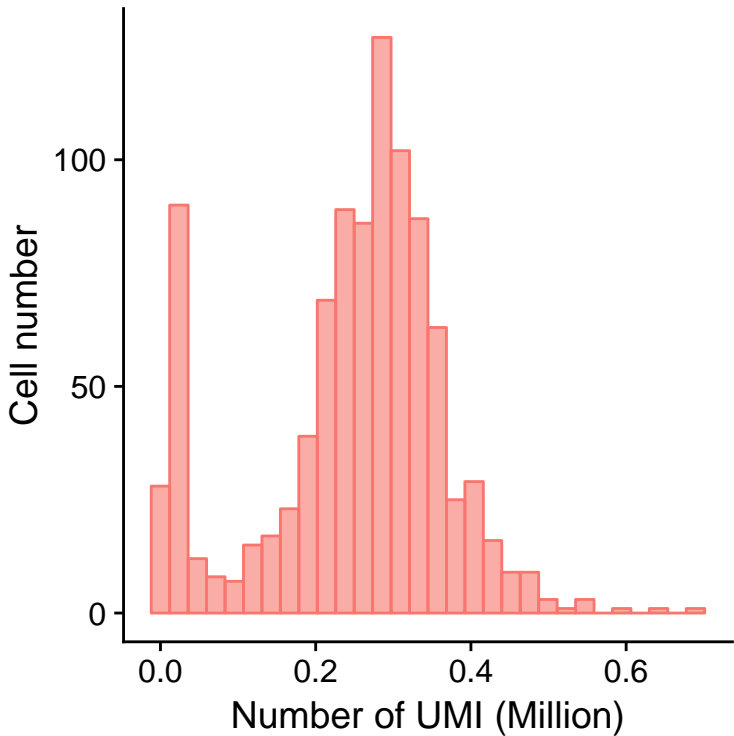
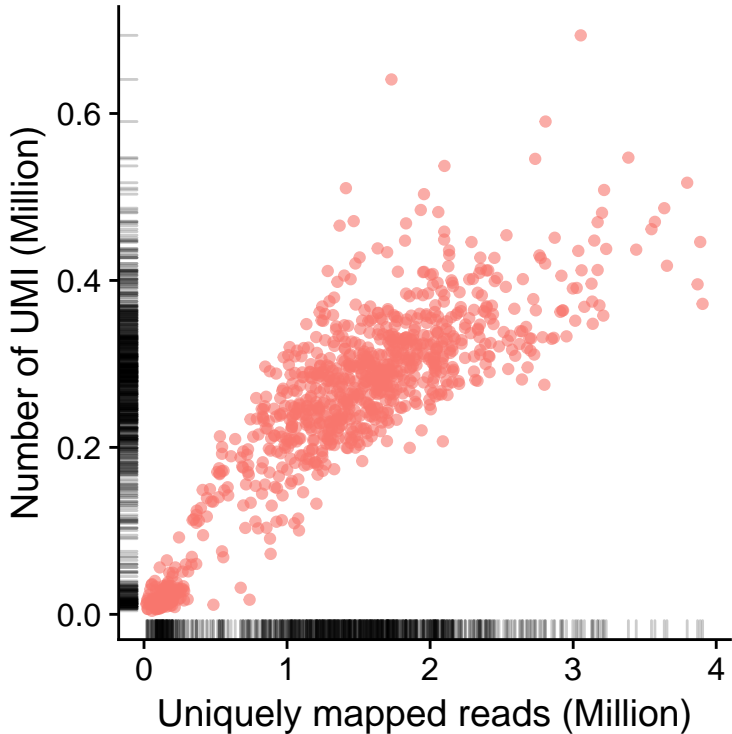


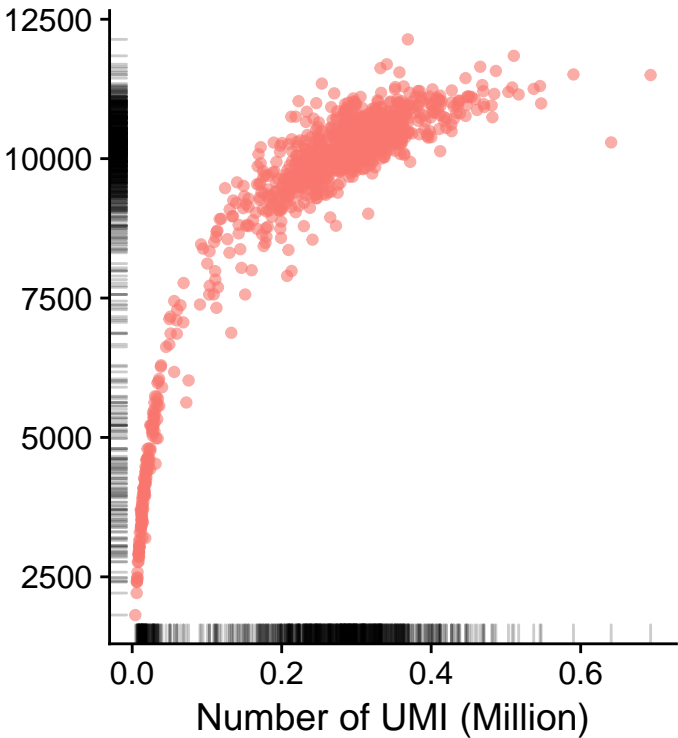
Number of detected genes

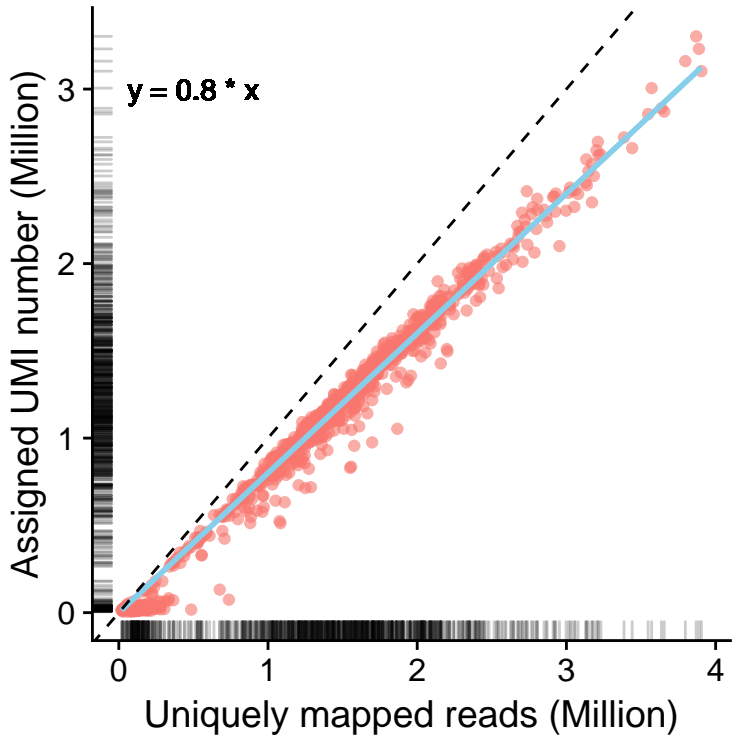


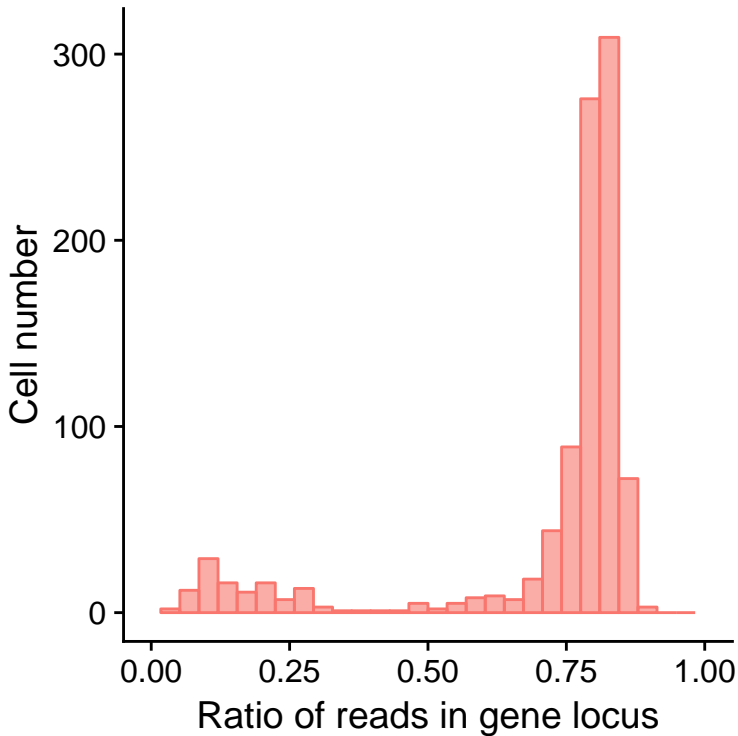


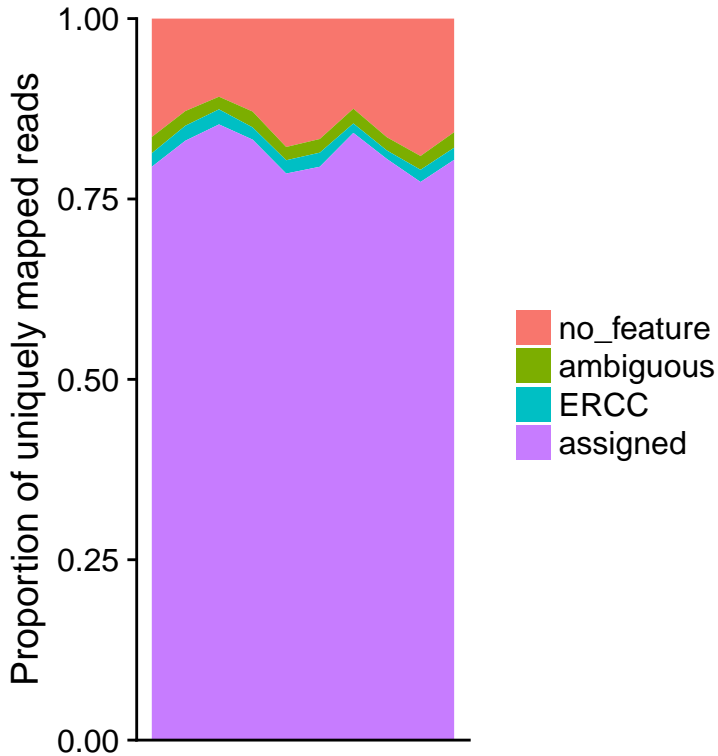


Number of detected genes

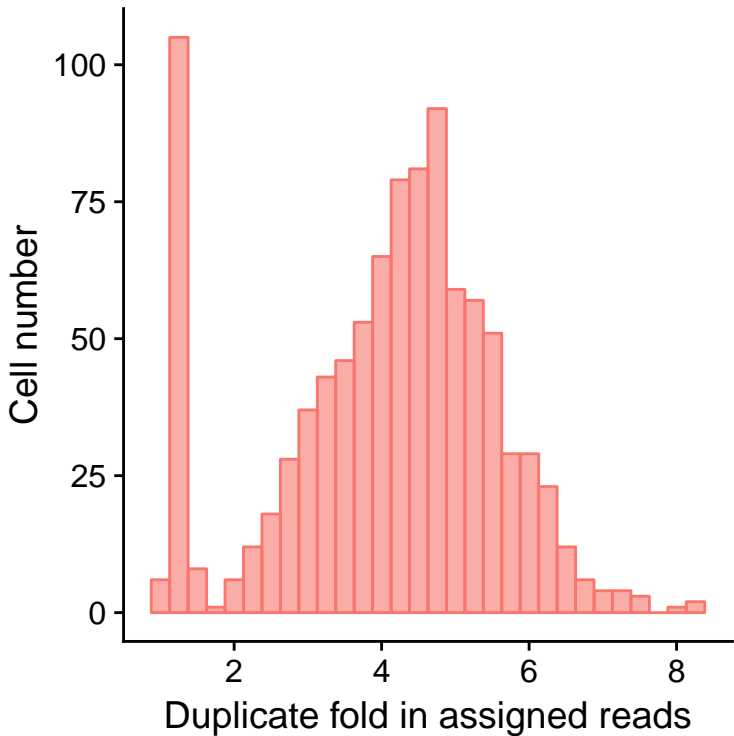












Duplicate fold in ERCC reads

12

8

4

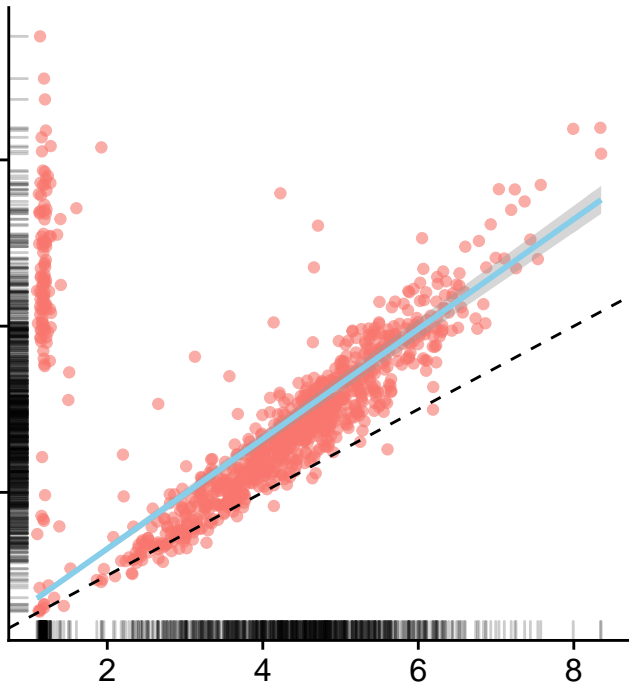
2

4

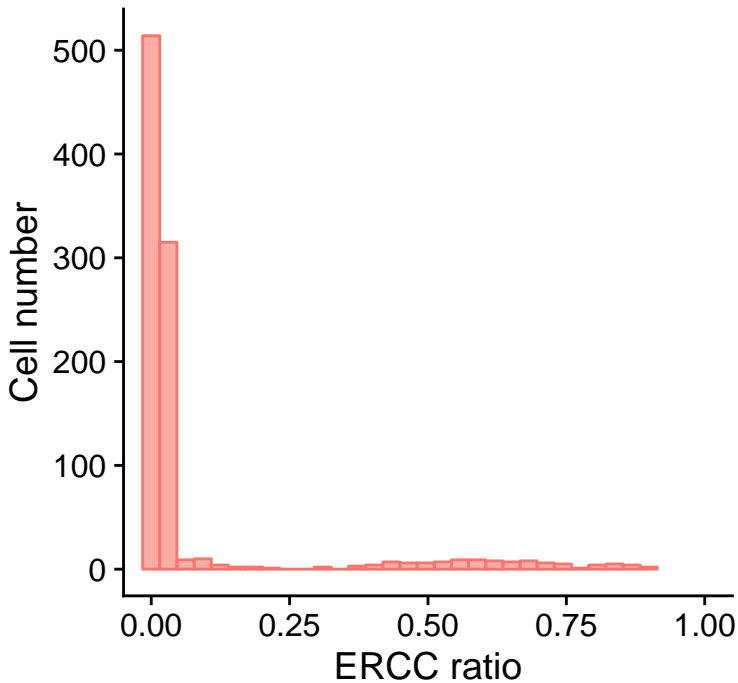
6

8

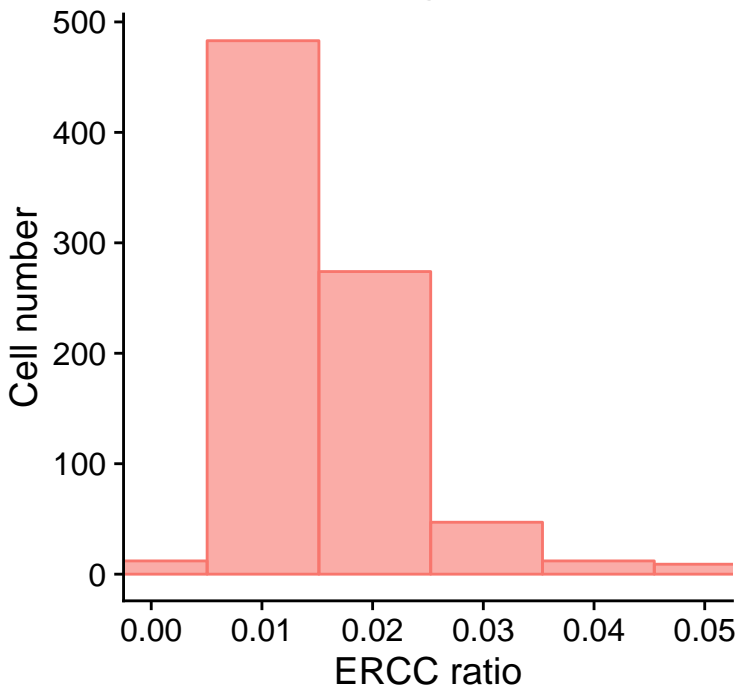
Duplicate fold in assigned reads



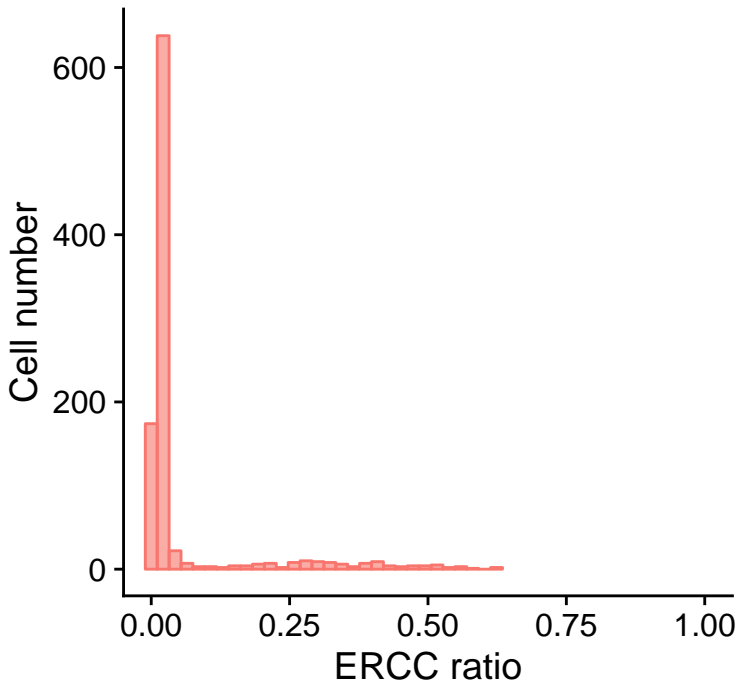
## Before removing duplicates



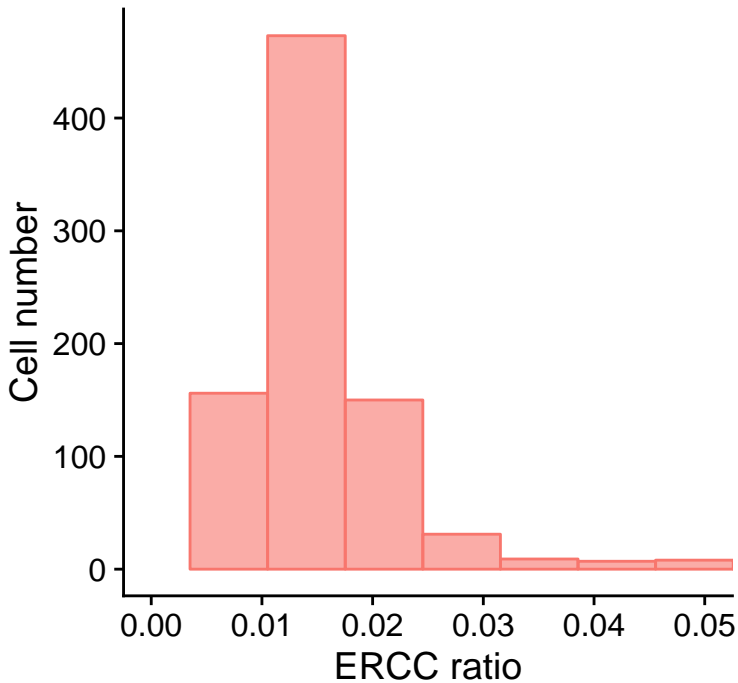
**Before removing dup. (Zoom in)**



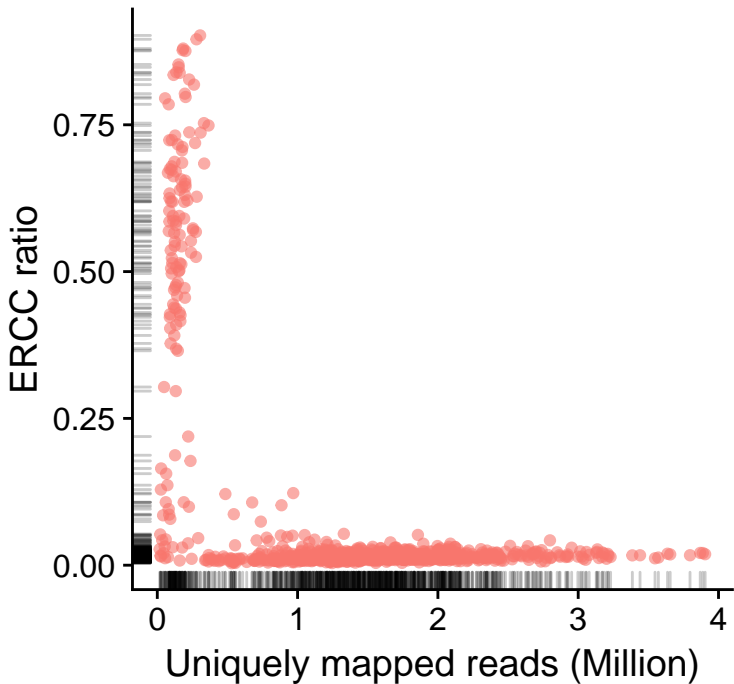
## After removing duplicates



## After removing dup. (Zoom in)



## Before removing duplicates



## After removing duplicates

