

Number of detected genes

12500  
10000  
7500  
5000  
2500  
0

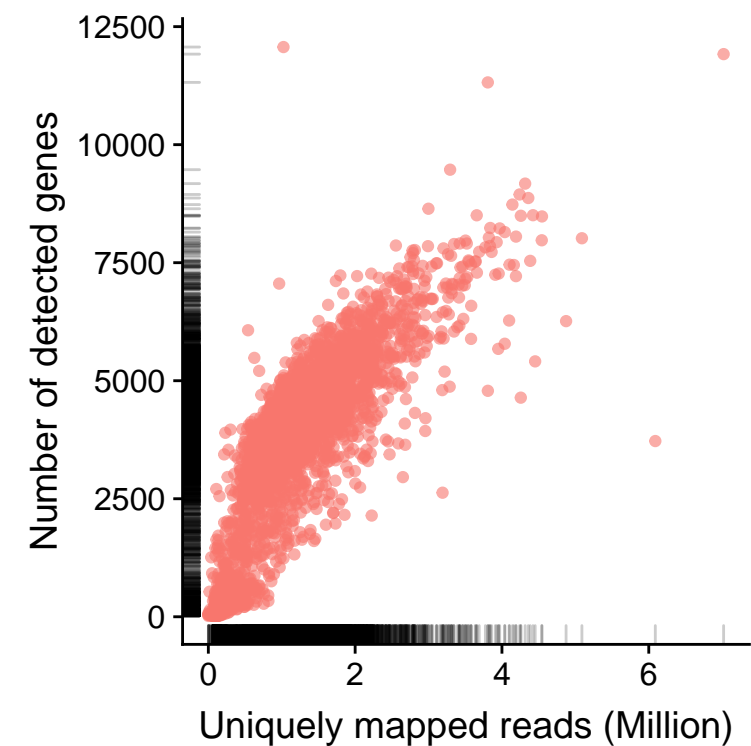
Uniquely mapped reads (Million)

0

2

4

6



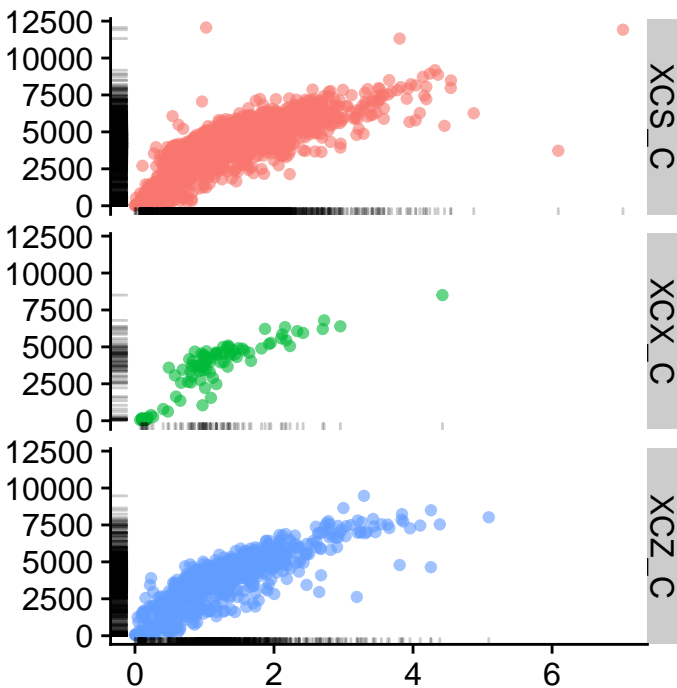
Number of detected genes

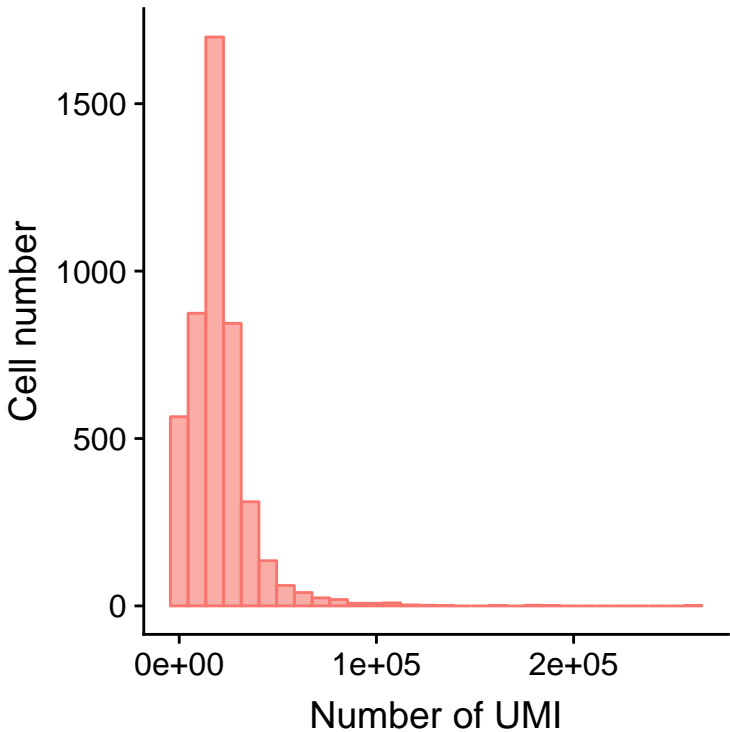
XCS\_C

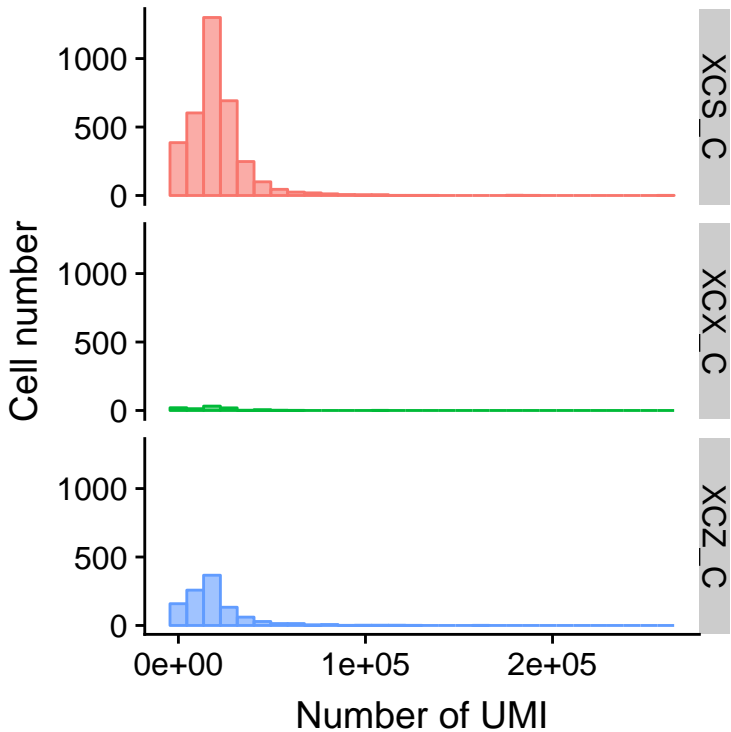
XCX\_C

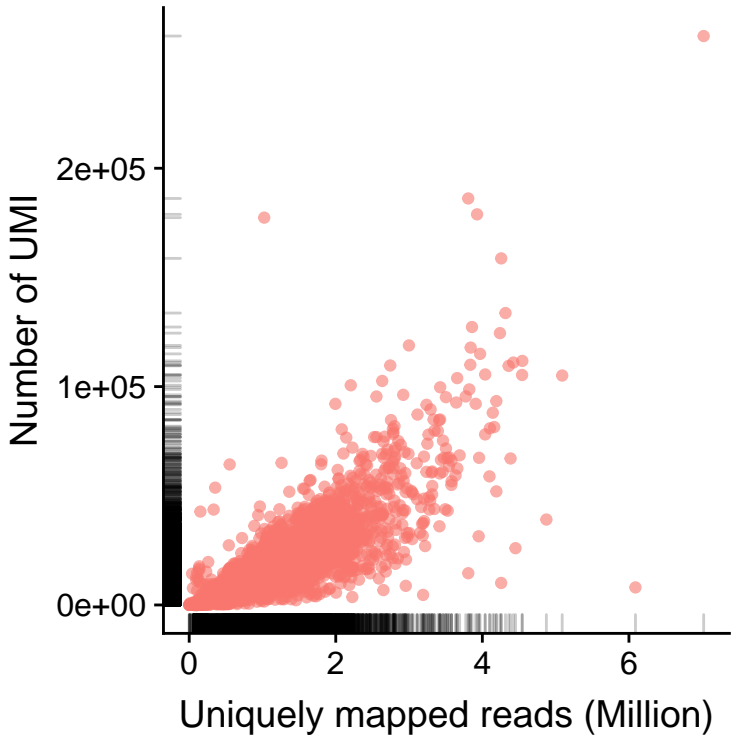
XCZ\_C

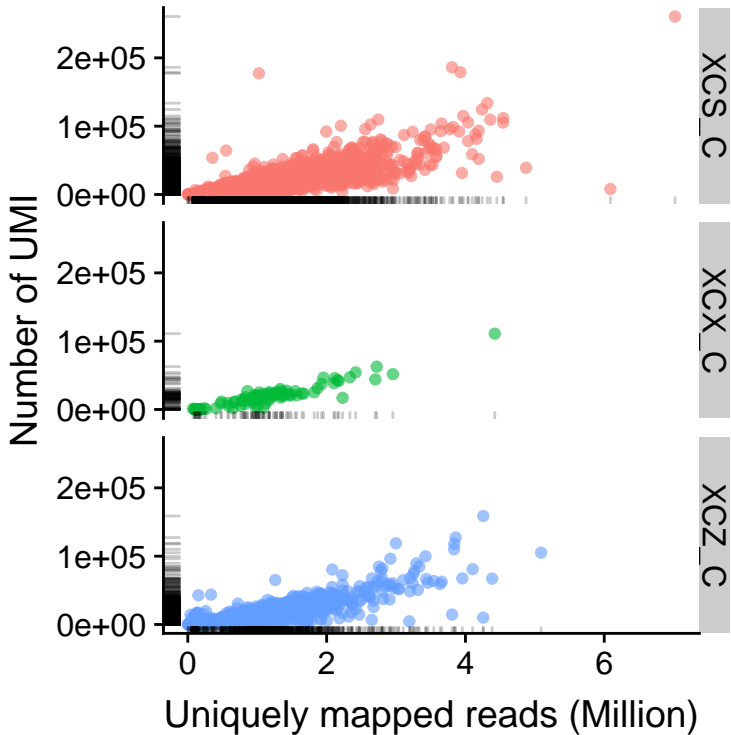
Uniquely mapped reads (Million)





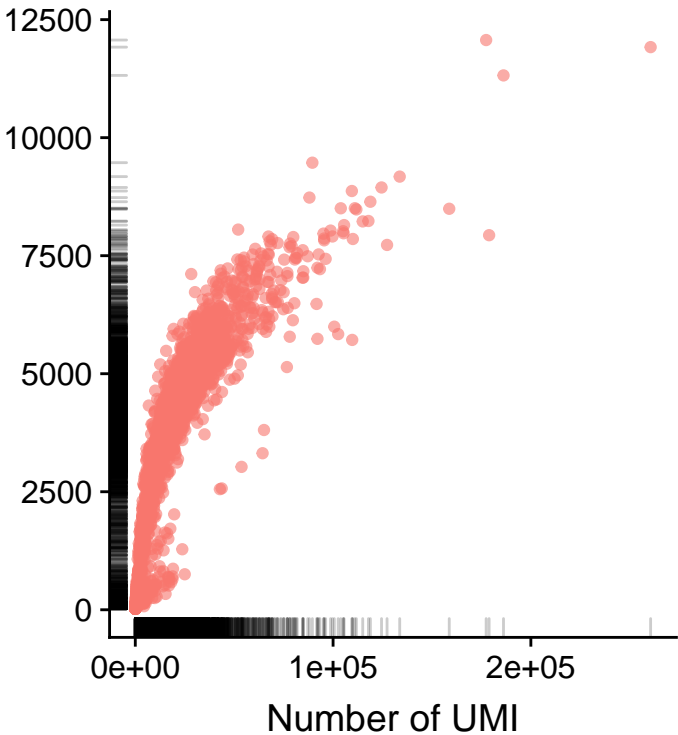


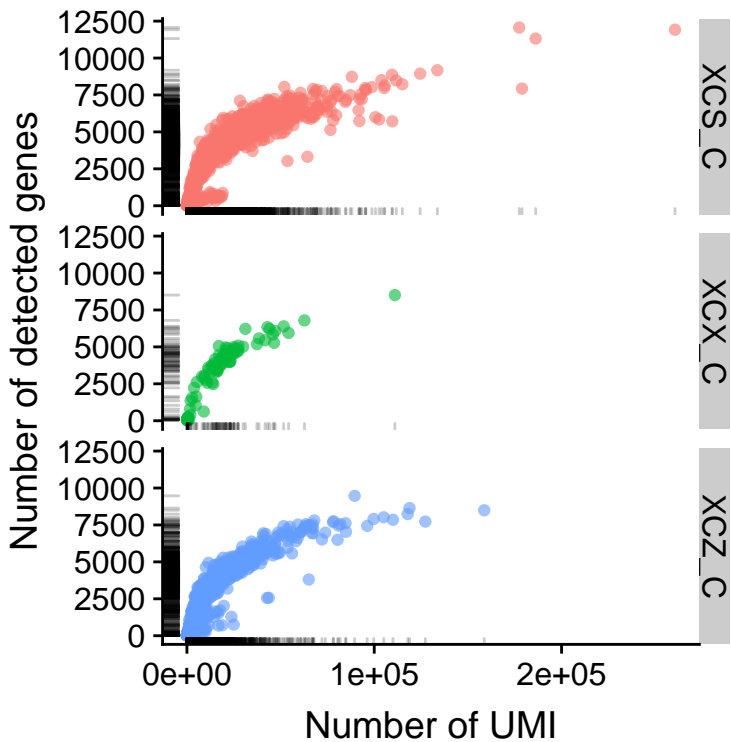


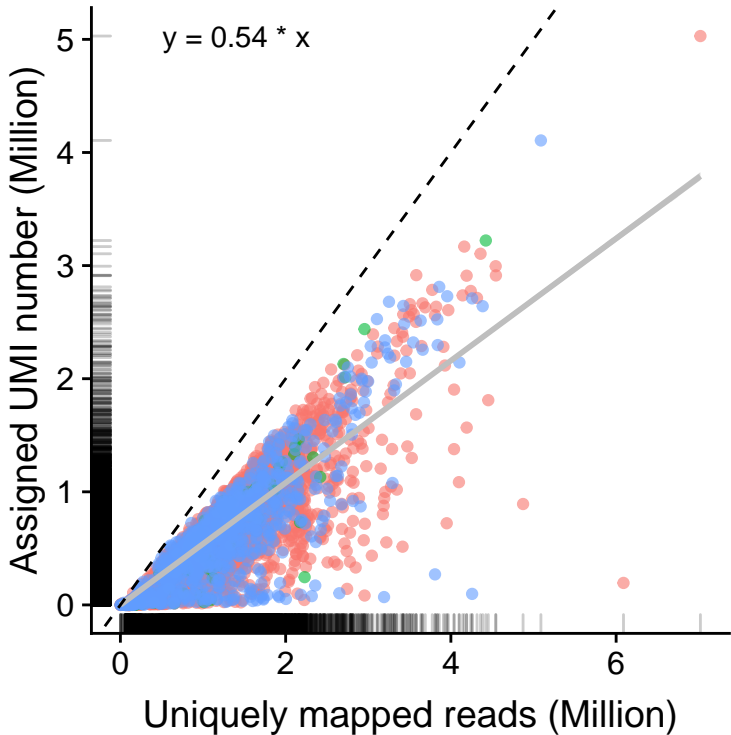


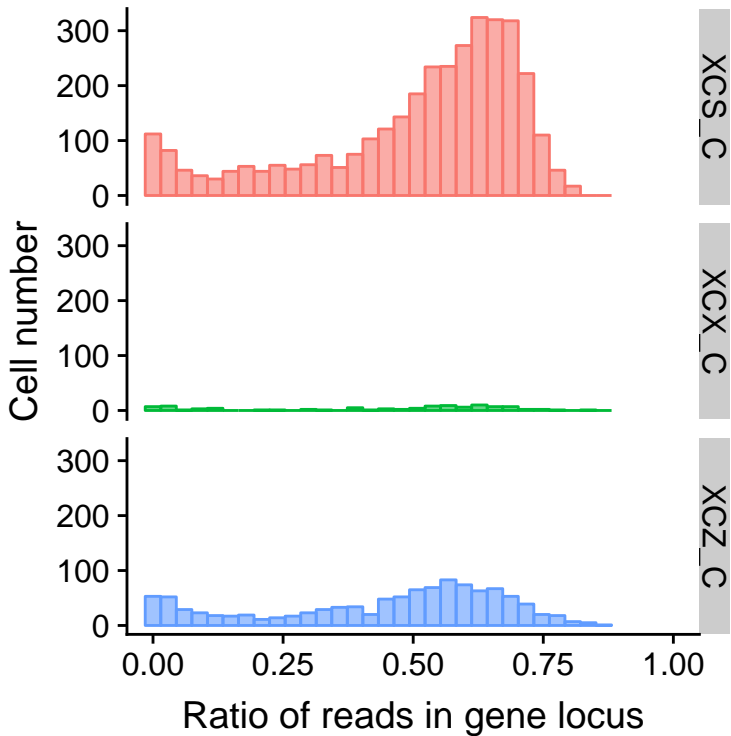


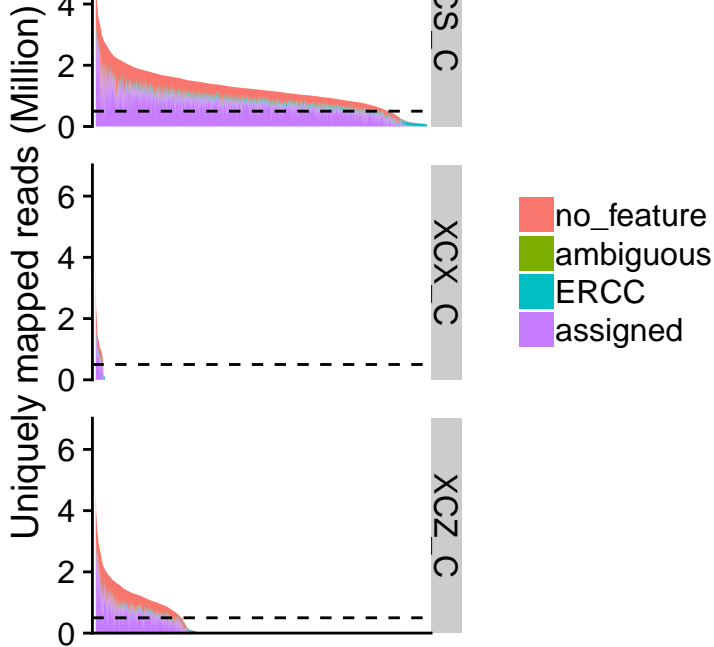
Number of detected genes

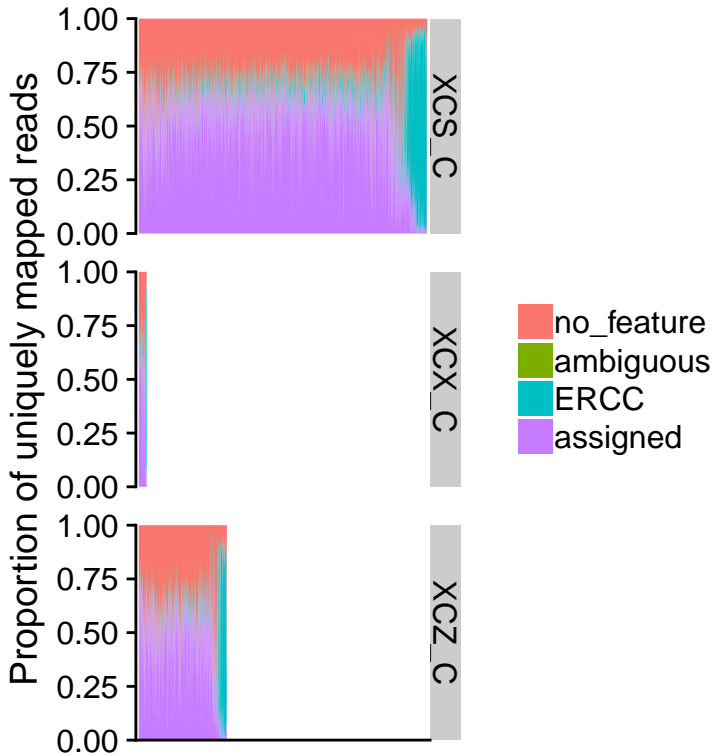


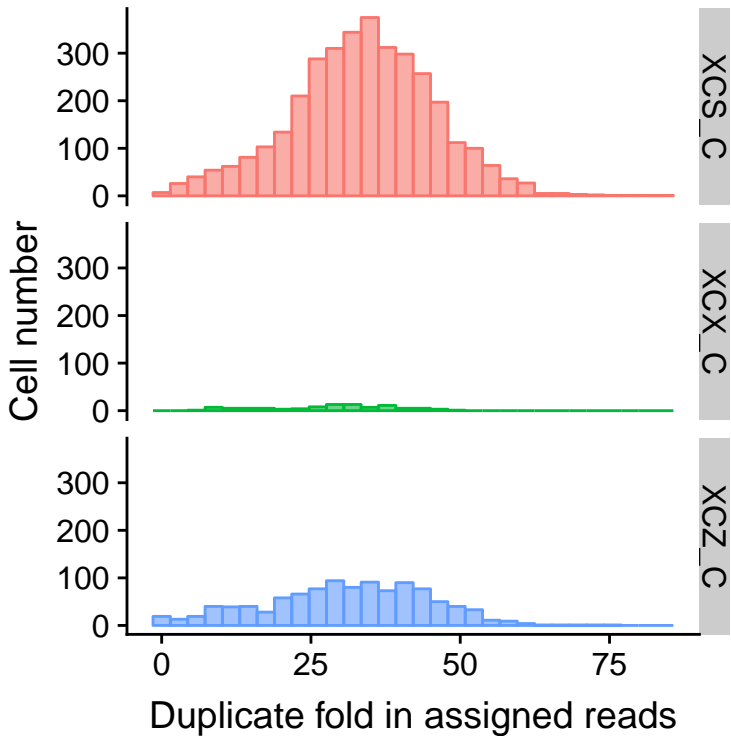




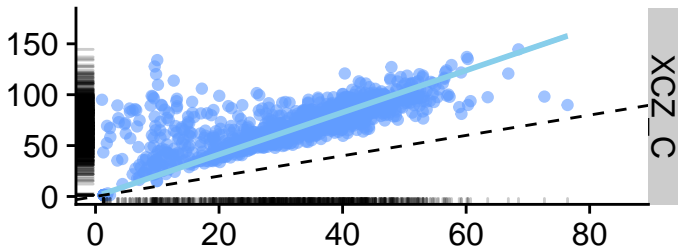
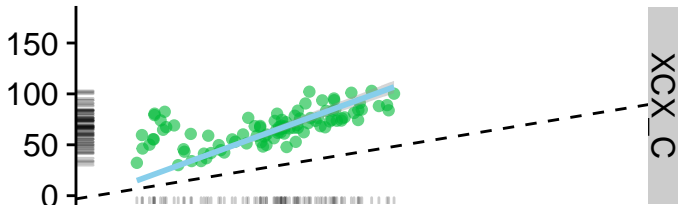
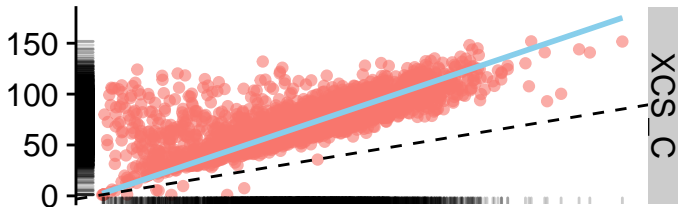








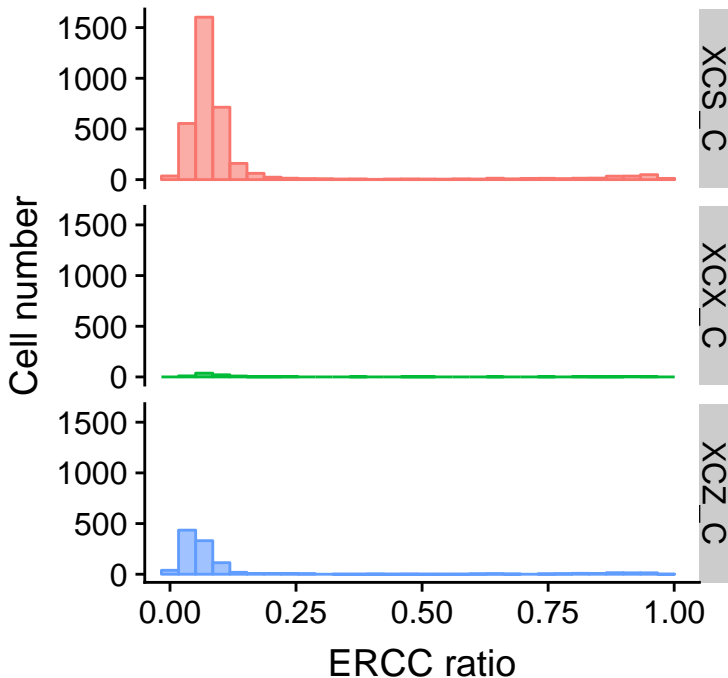
Duplicate fold in ERCC reads



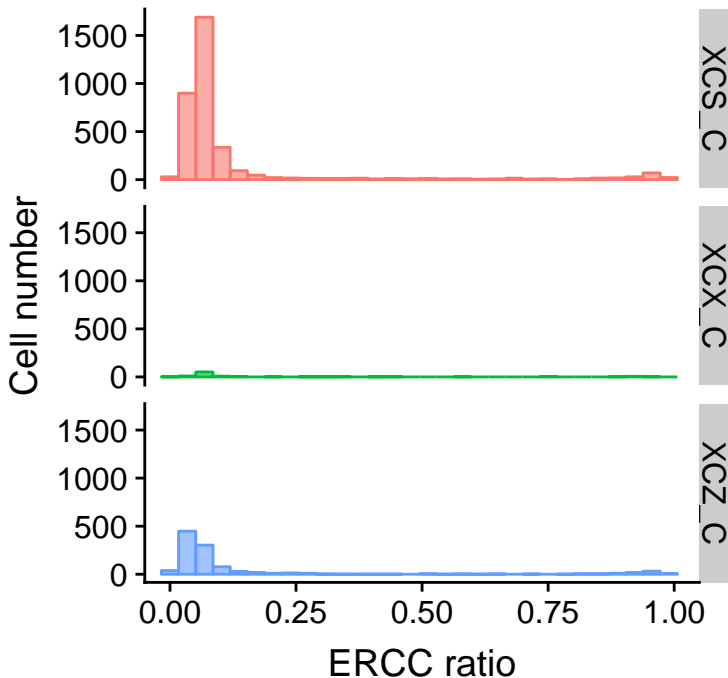
Duplicate fold in assigned reads



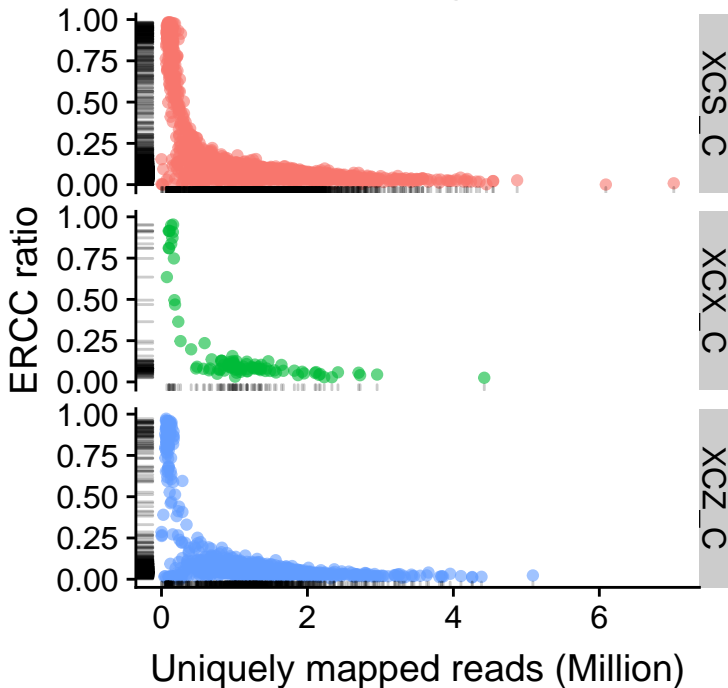
## Before removing duplicates



## After removing duplicates



## Before removing duplicates



## After removing duplicates

