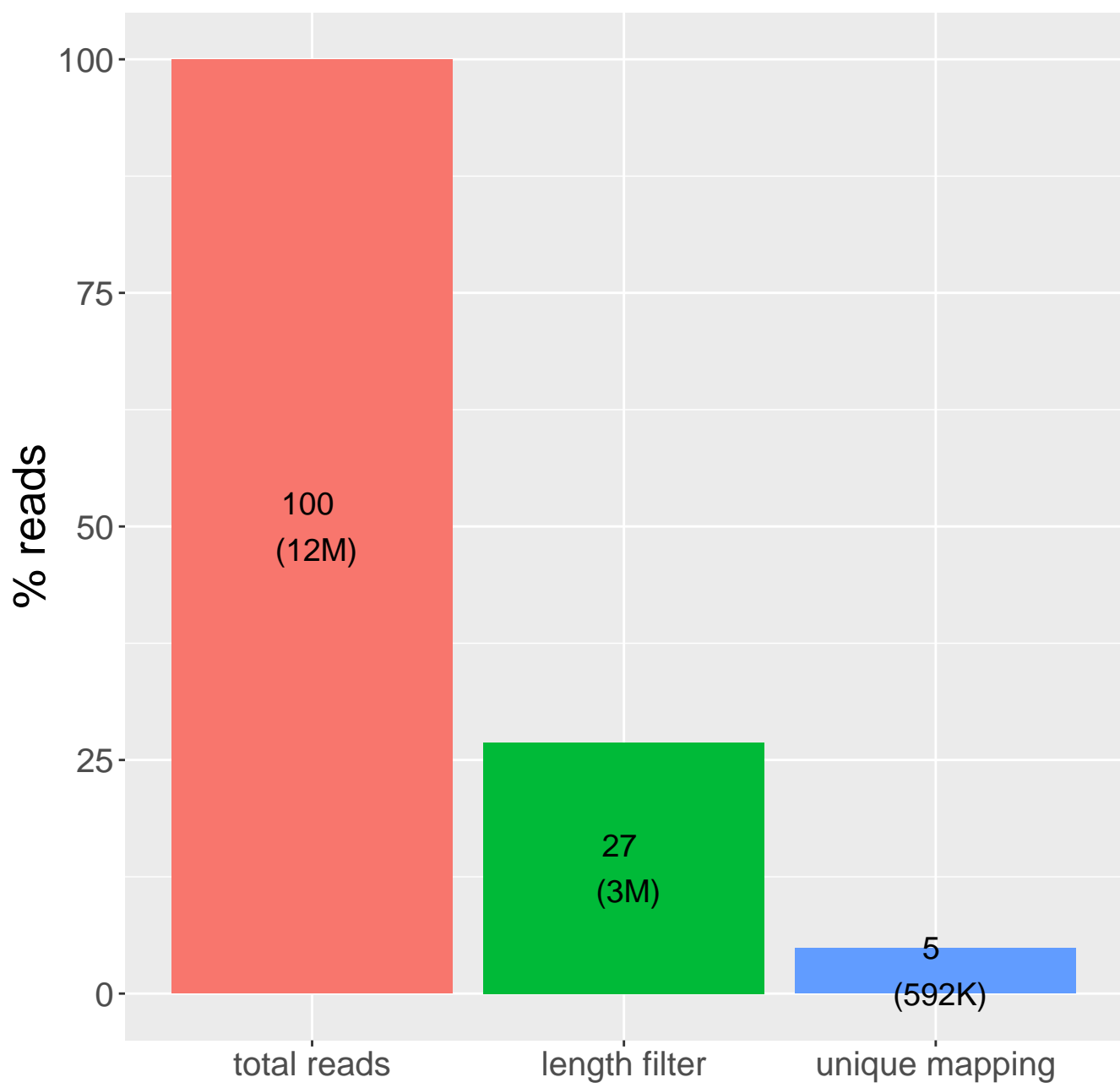
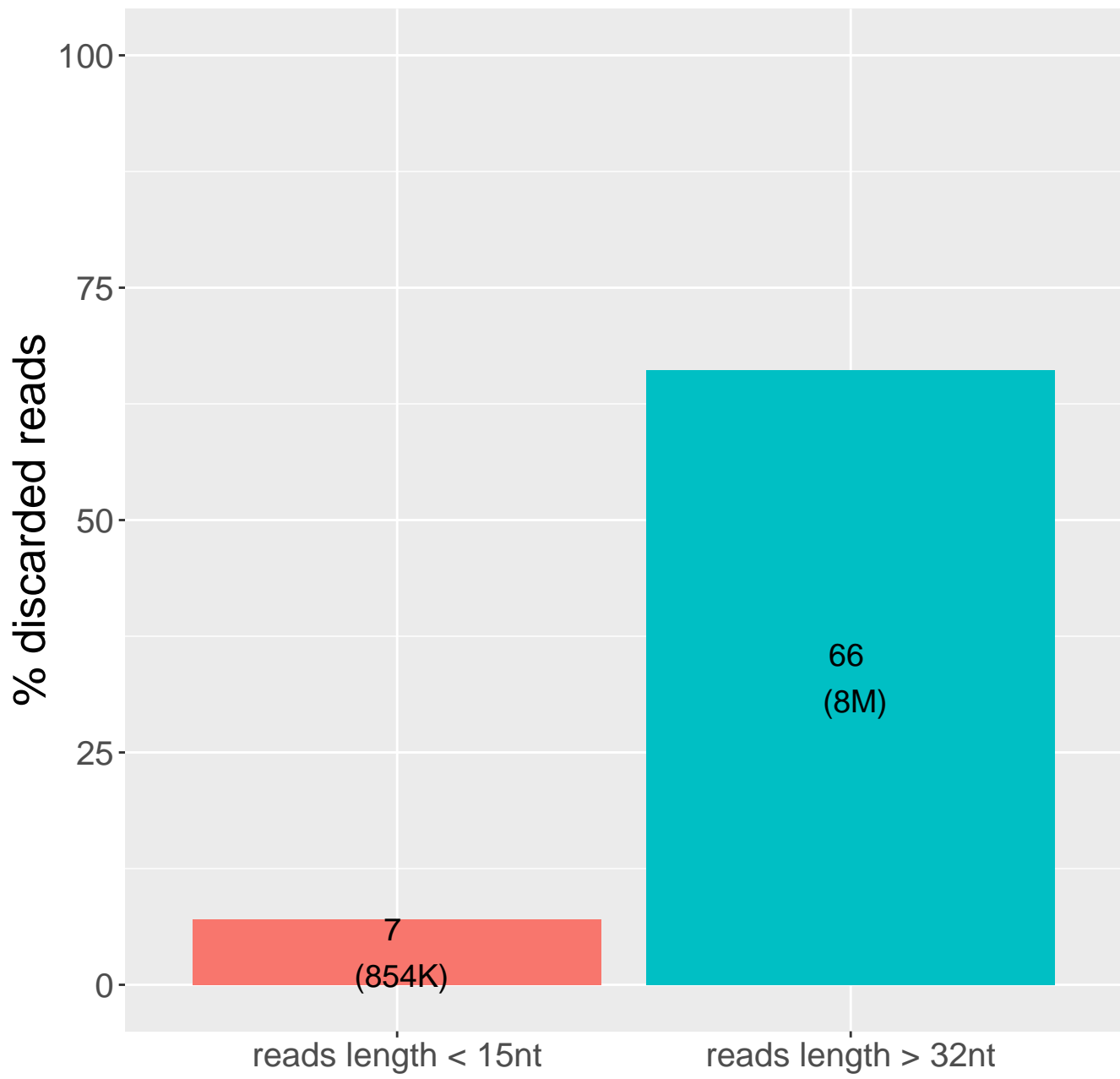


Retained reads

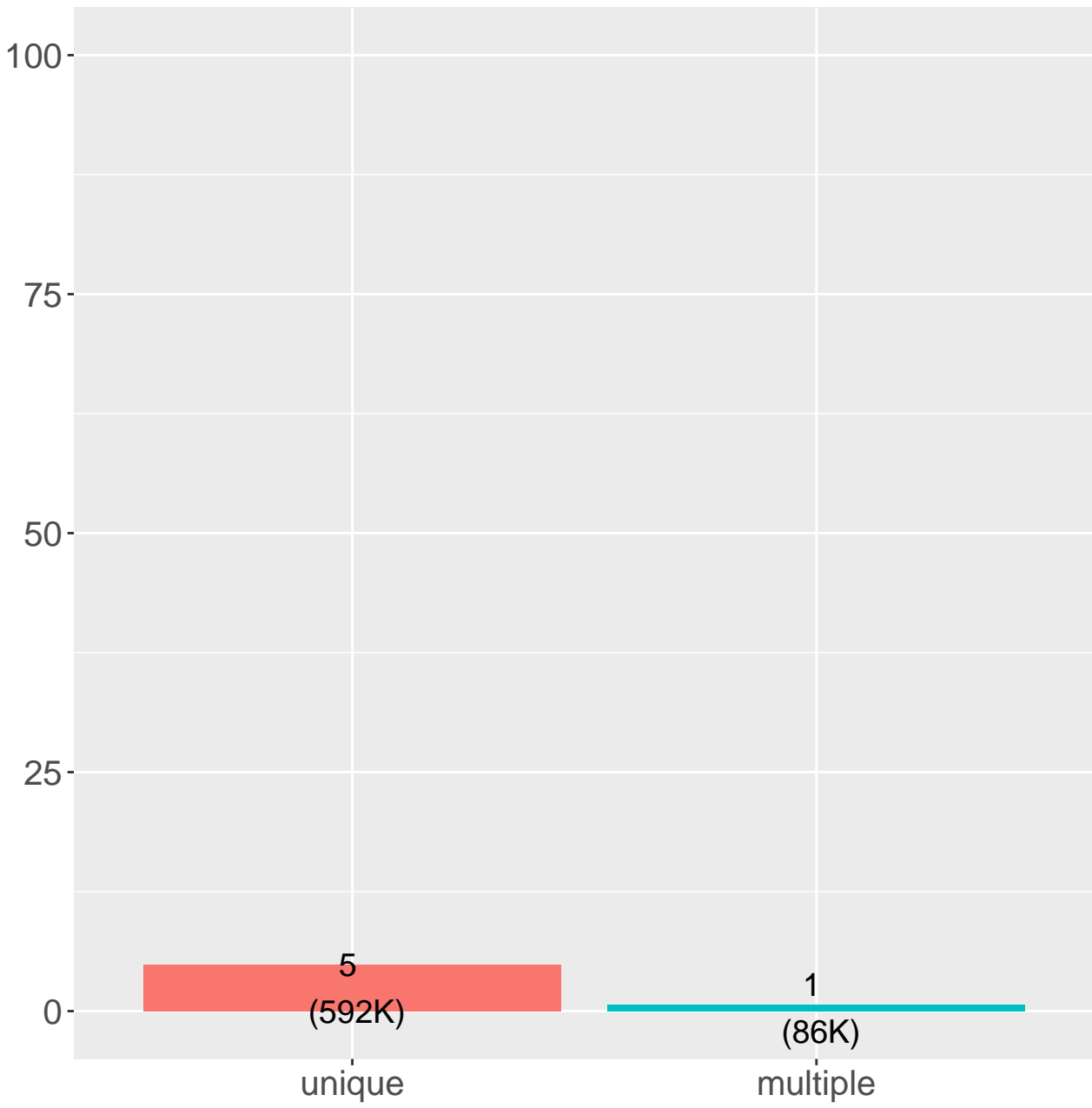


Length filtering



Mapping reads

% mapped reads



Read distribution

% reads

100

75

50

25

0

miRNA

piRNA

snoRNA

snRNA

rRNA

53
(312K)

26
(156K)

3
(19K)

1
(7K)

4
(25K)

