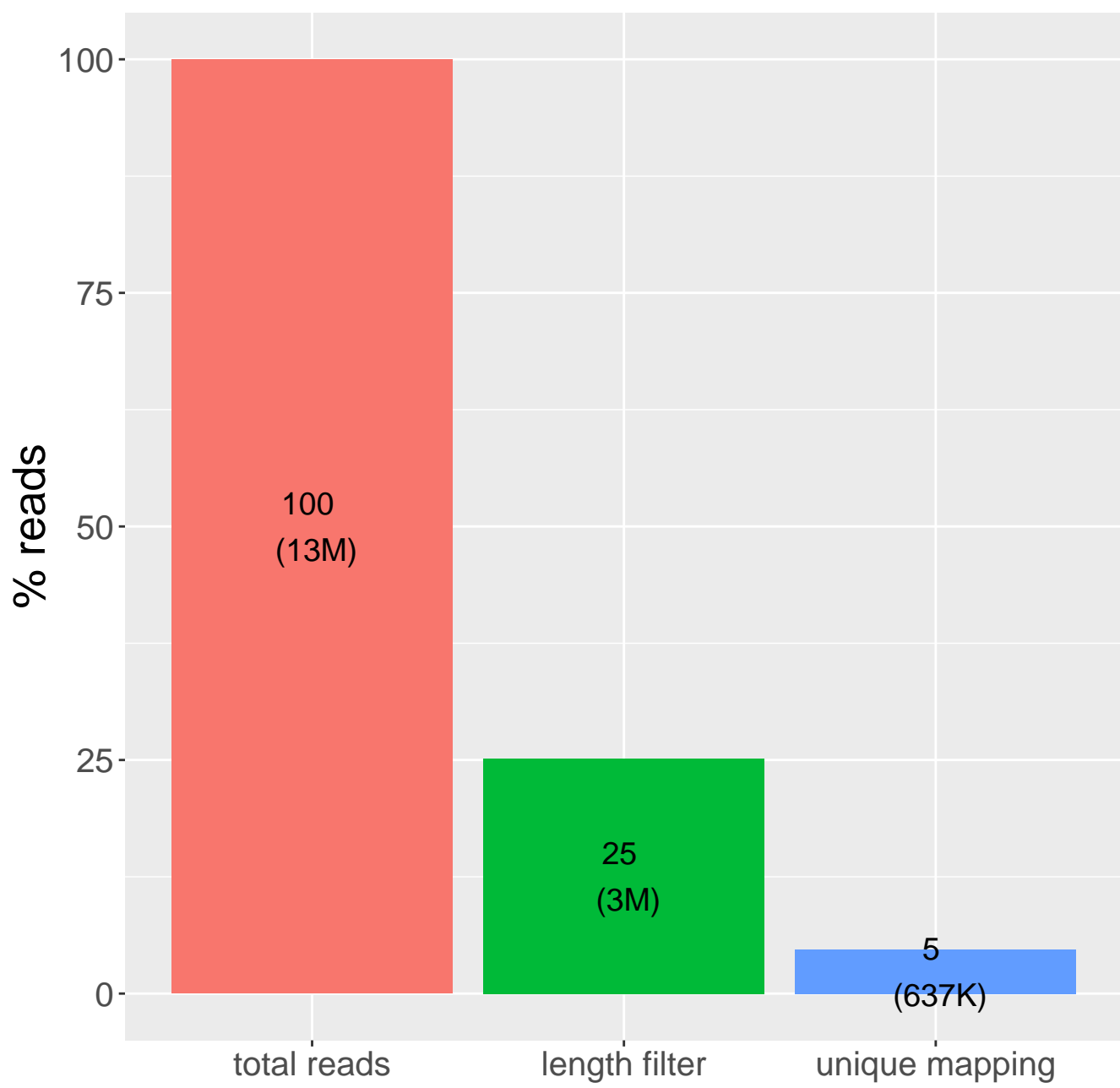
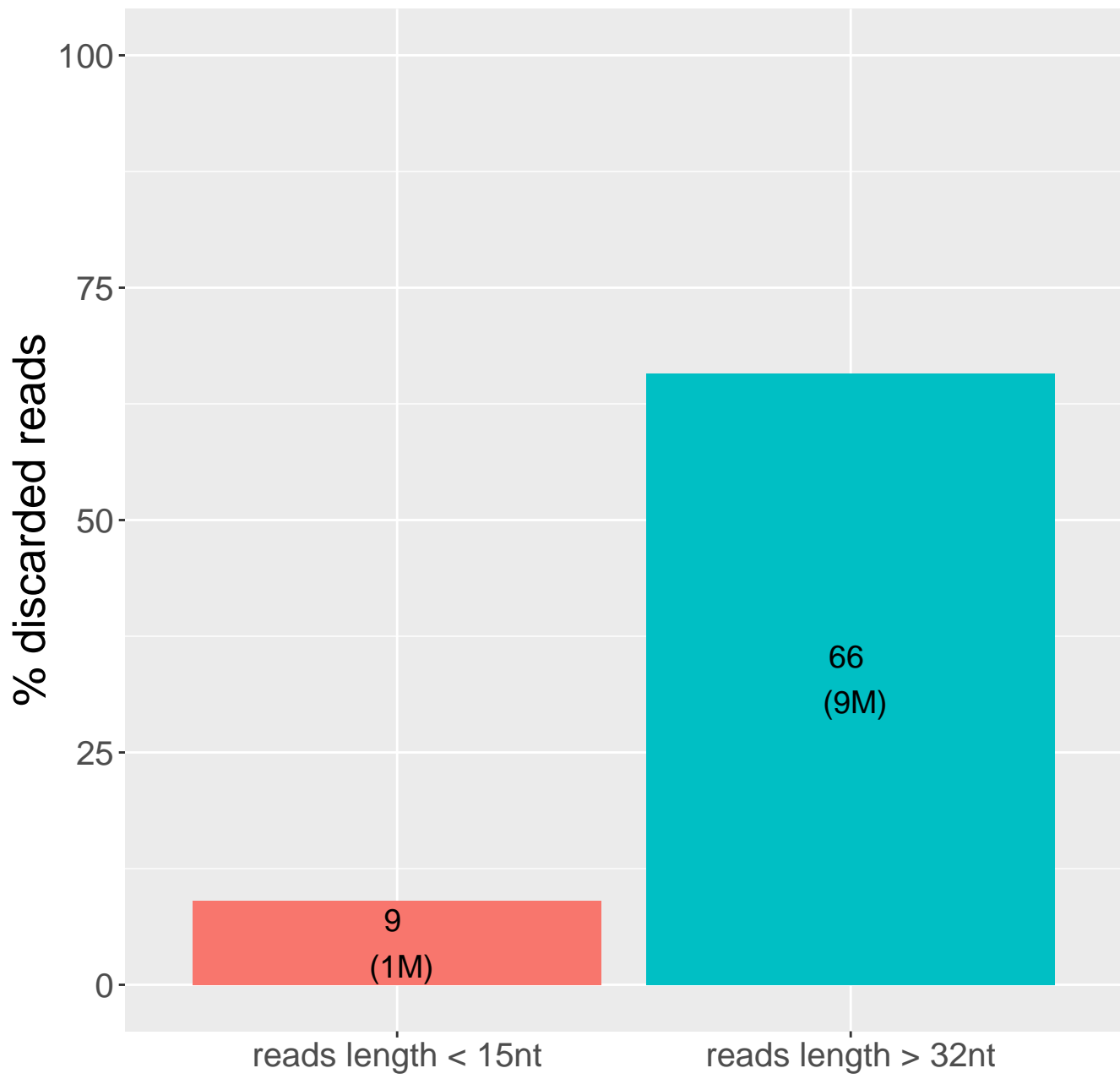


Retained reads

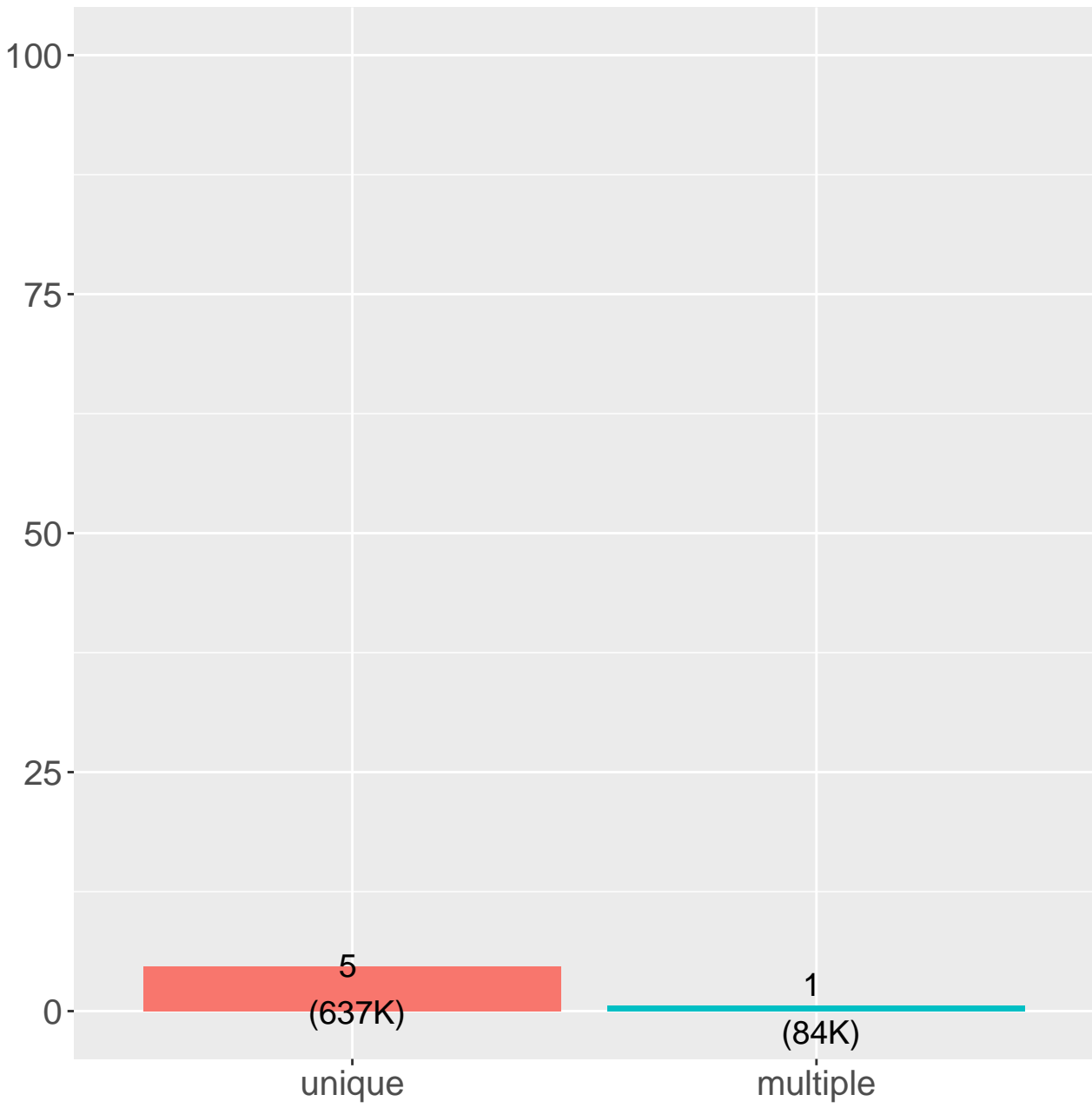


Length filtering



Mapping reads

% mapped reads



Read distribution

% reads

100

75

50

25

0

miRNA

piRNA

snoRNA

snRNA

rRNA

59
(374K)

26
(165K)

3
(18K)

1
(4K)

4
(27K)

