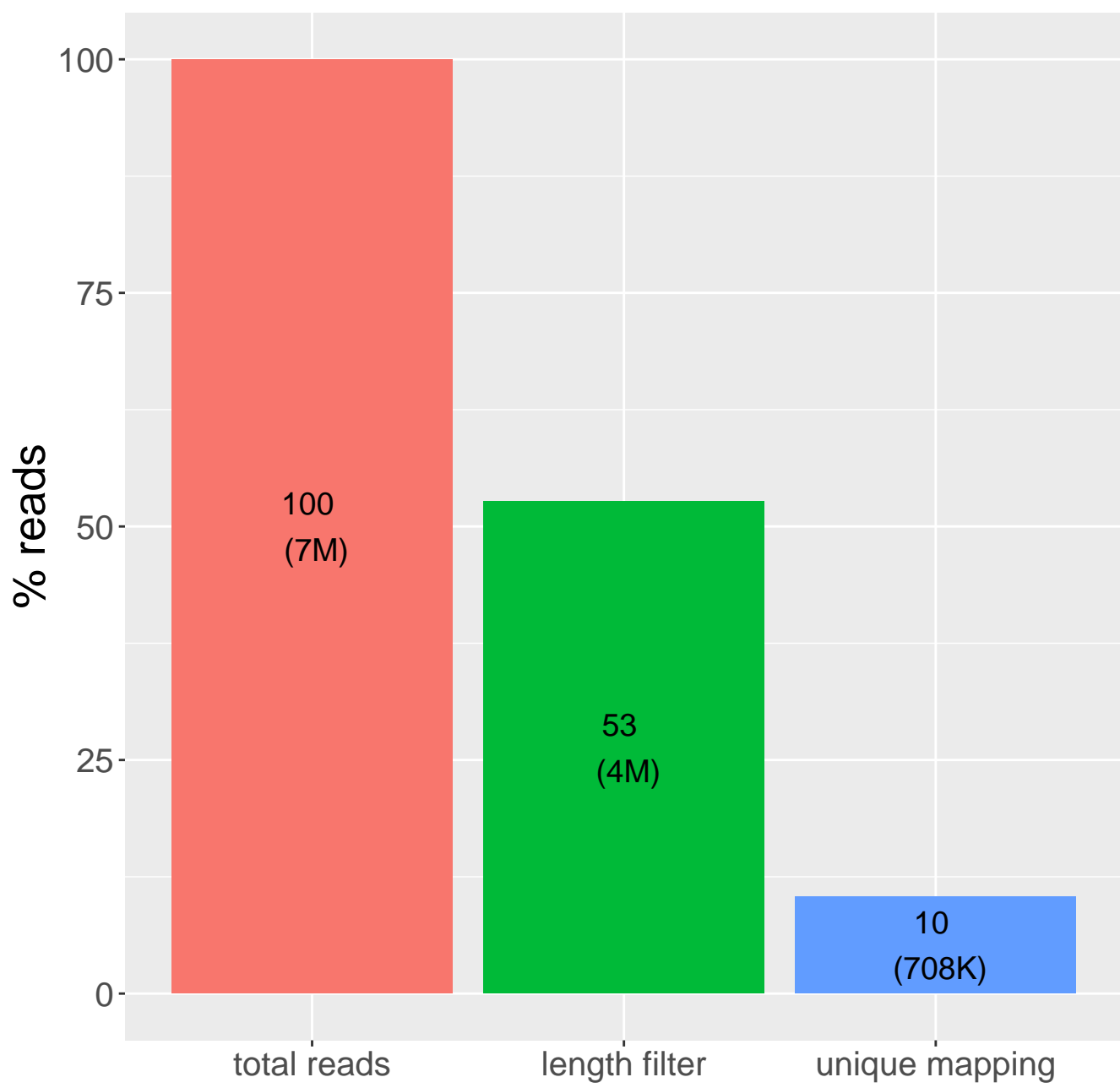
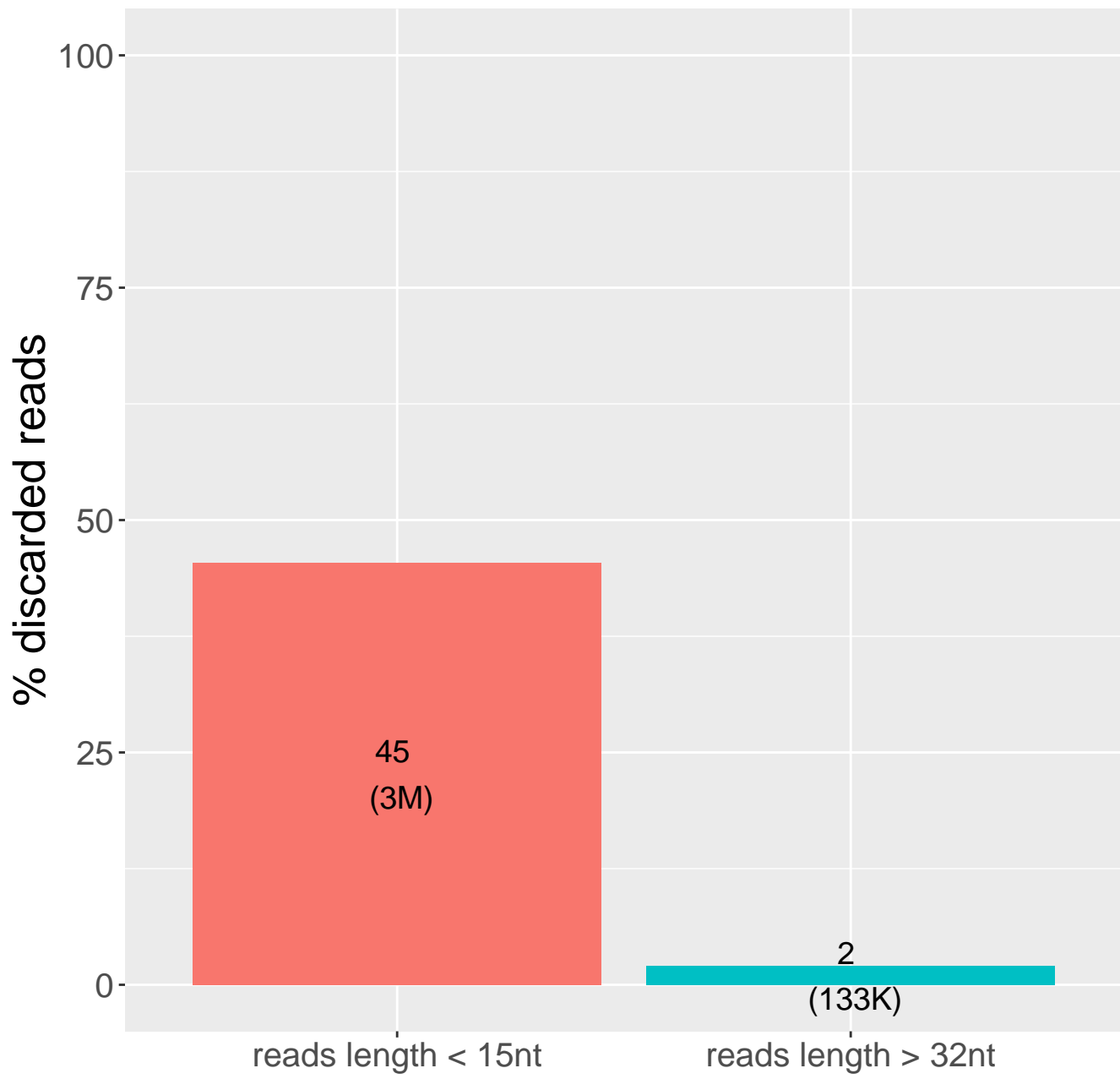


# Retained reads

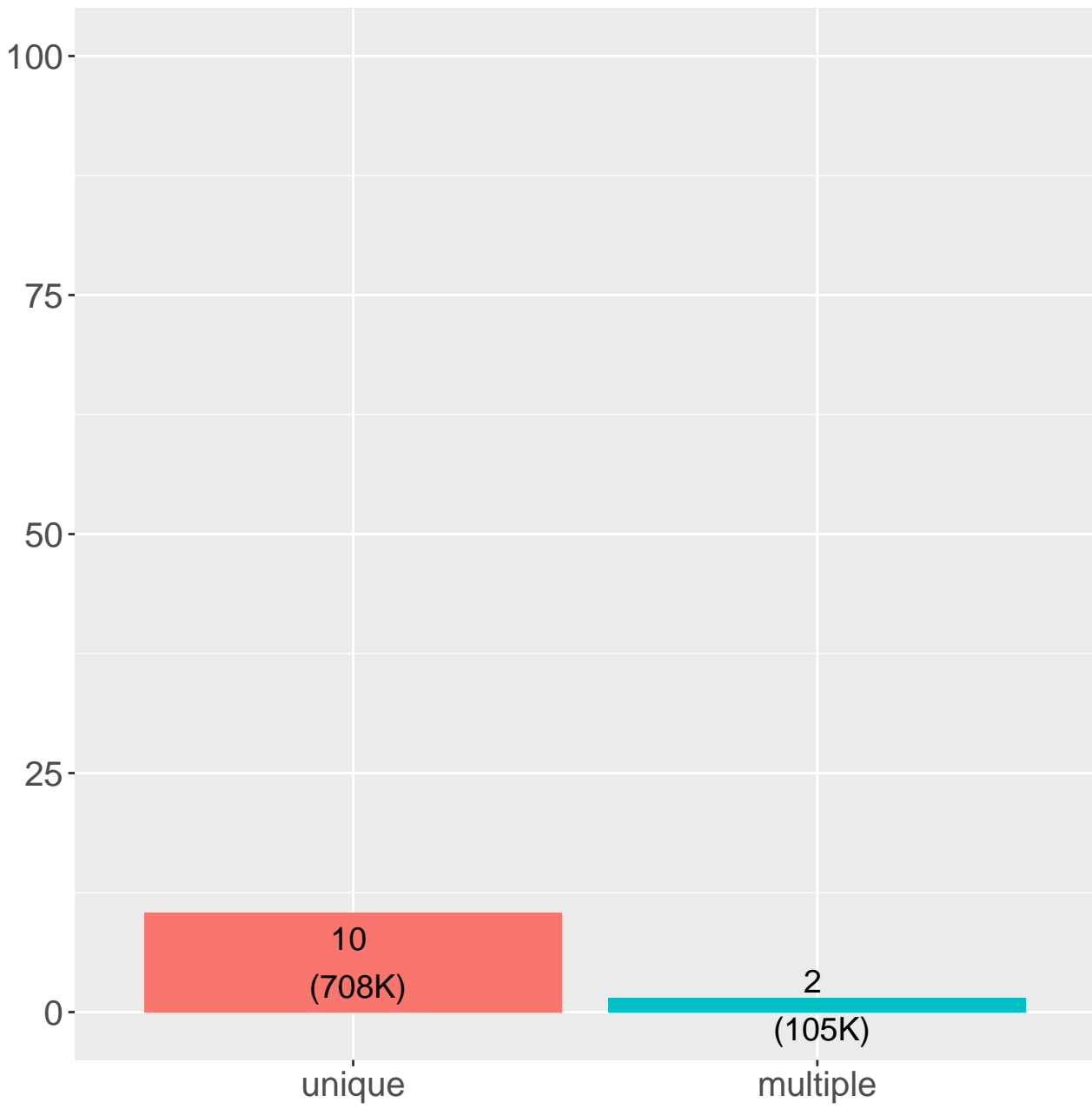


# Length filtering



# Mapping reads

% mapped reads



# Read distribution

% reads

100

75

50

25

0

miRNA

piRNA

snoRNA

snRNA

rRNA

66  
(465K)

22  
(155K)

3  
(18K)

0  
(3K)

2  
(12K)

