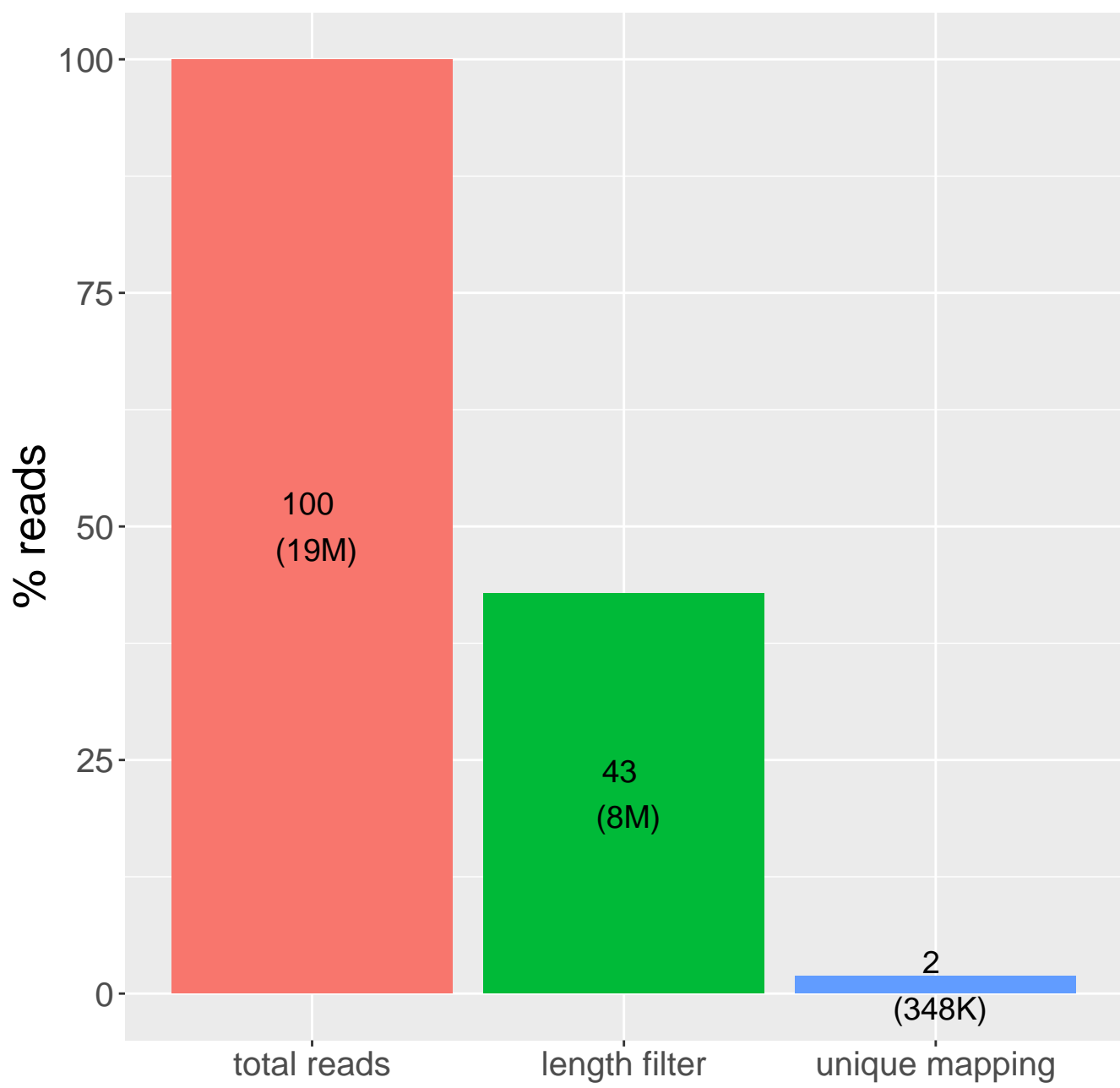
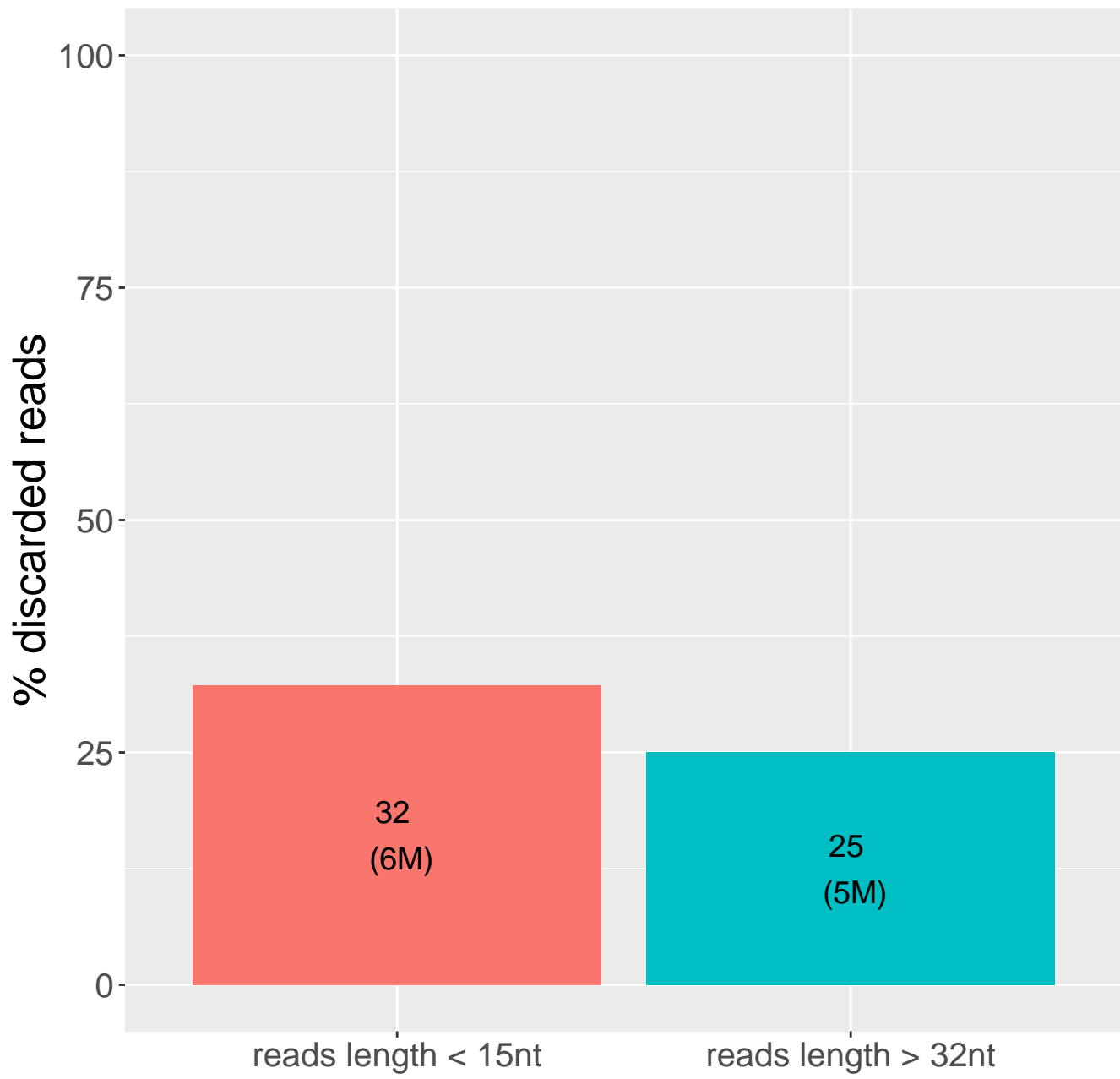


# Retained reads

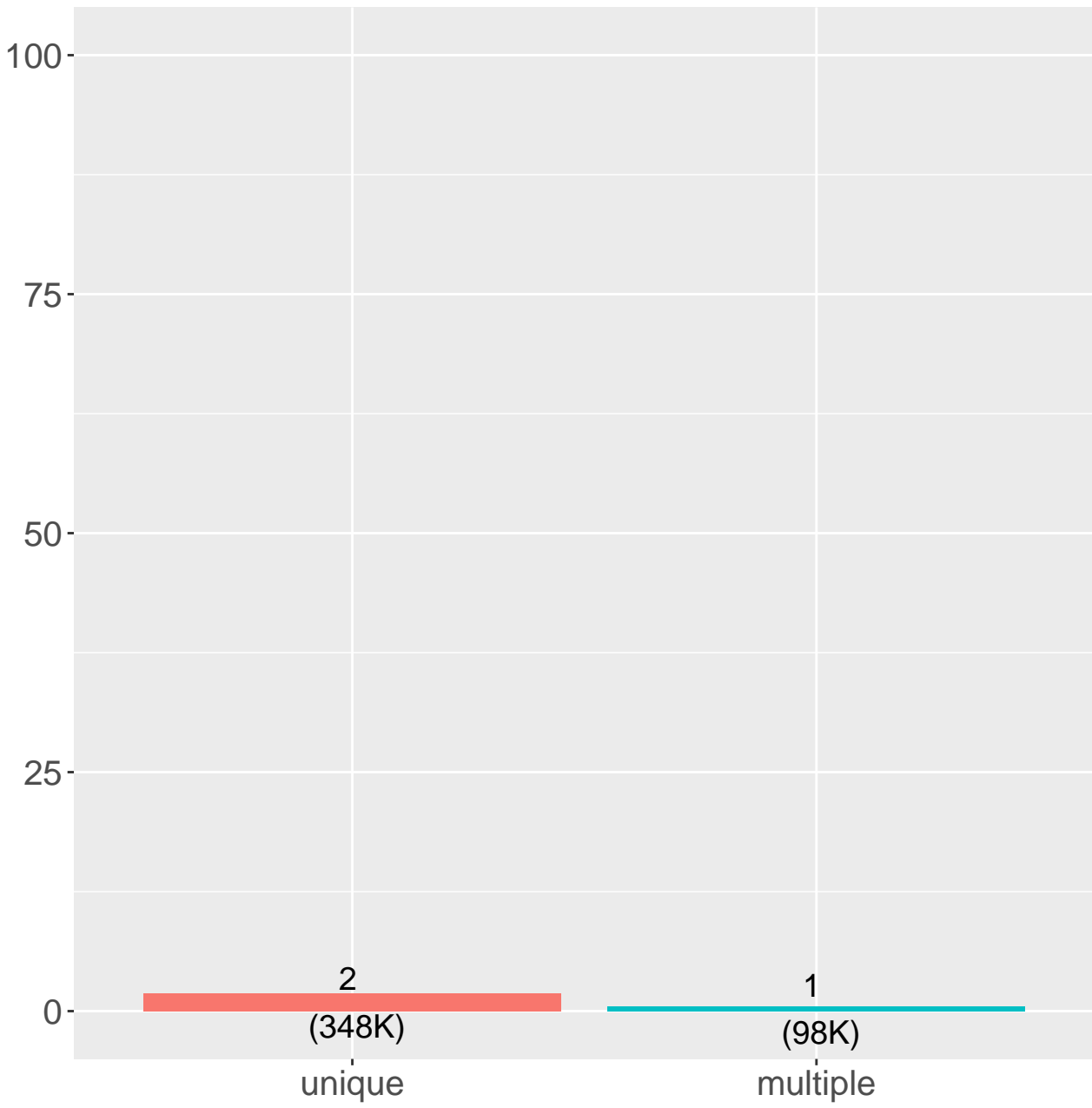


# Length filtering



# Mapping reads

% mapped reads



# Read distribution

% reads

100

75

50

25

0

miRNA

piRNA

snoRNA

snRNA

rRNA

36  
(126K)

45  
(156K)

2  
(6K)

1  
(4K)

4  
(13K)

