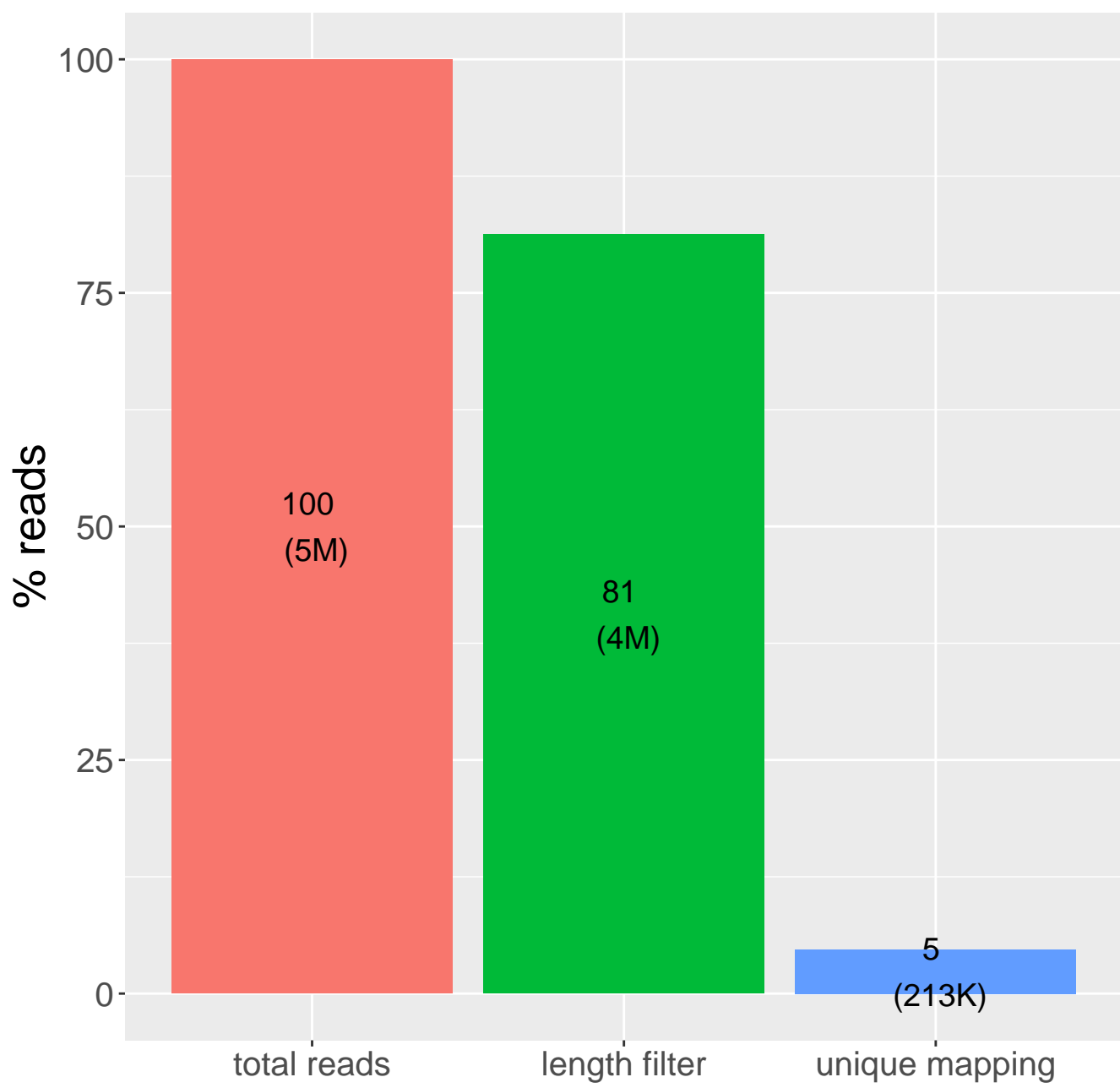
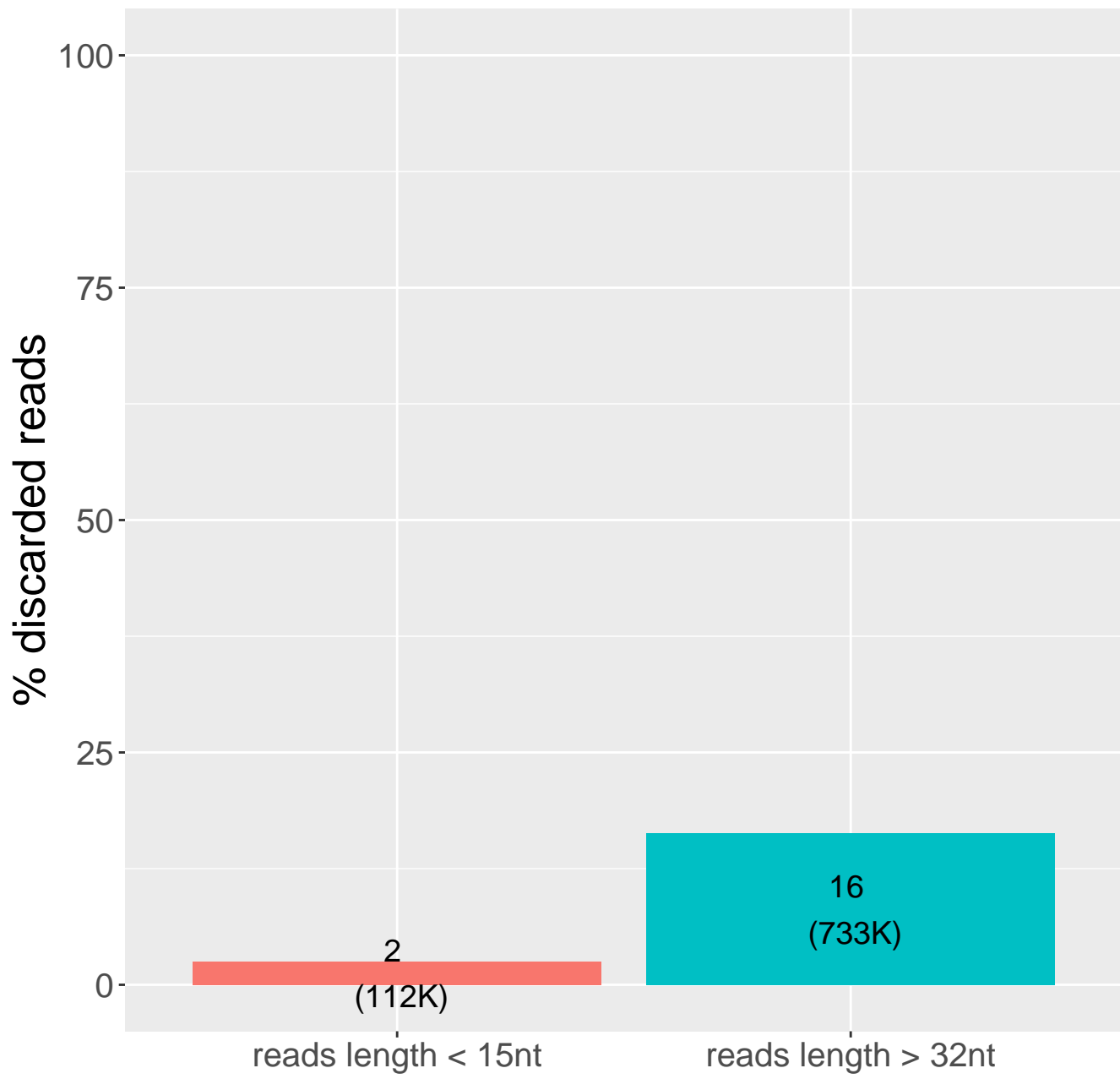


# Retained reads

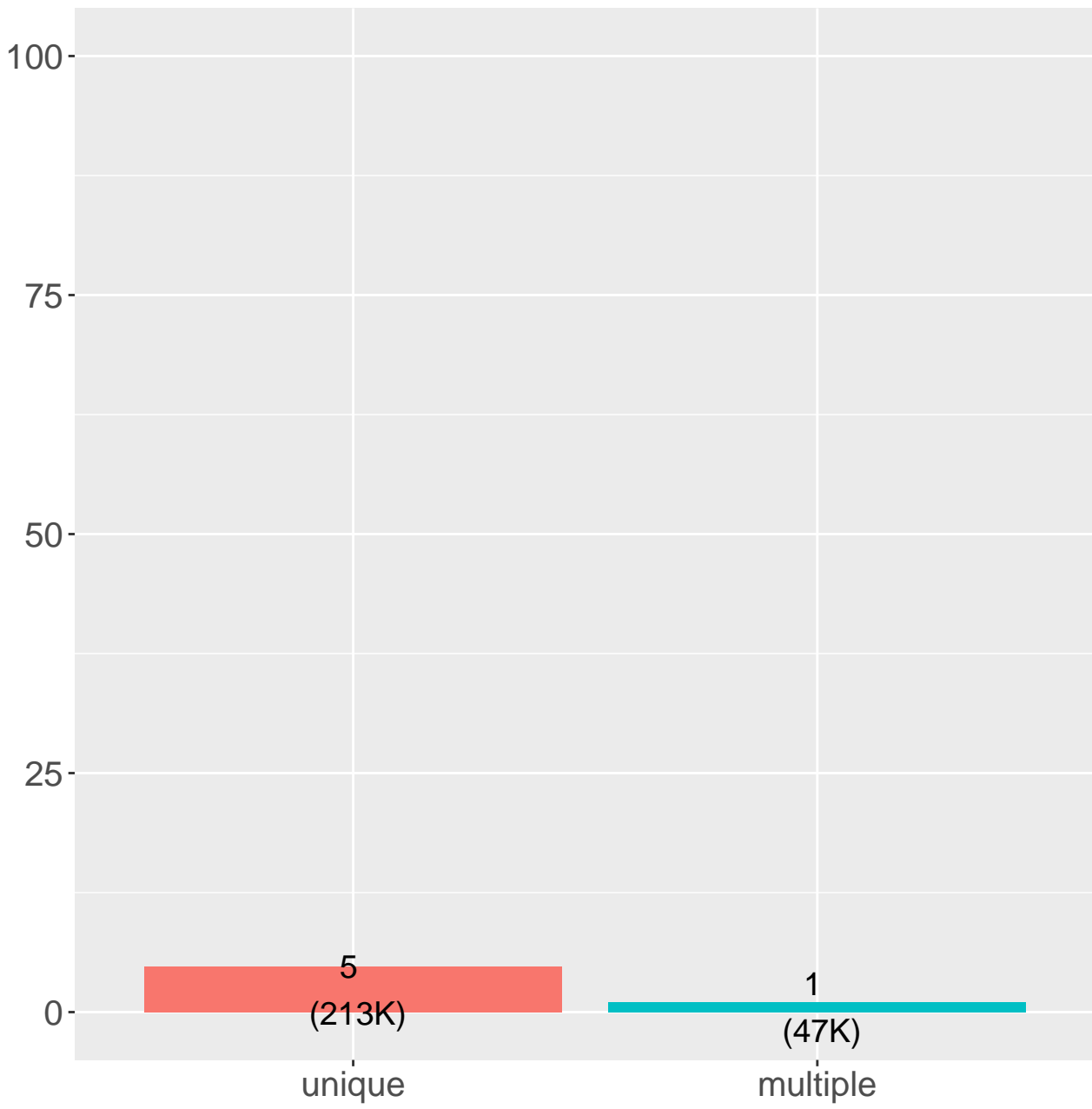


# Length filtering



# Mapping reads

% mapped reads



# Read distribution

% reads

100

75

50

25

0

miRNA

piRNA

snoRNA

snRNA

rRNA

51  
(109K)

29  
(61K)

10  
(21K)

1  
(2K)

4  
(8K)

