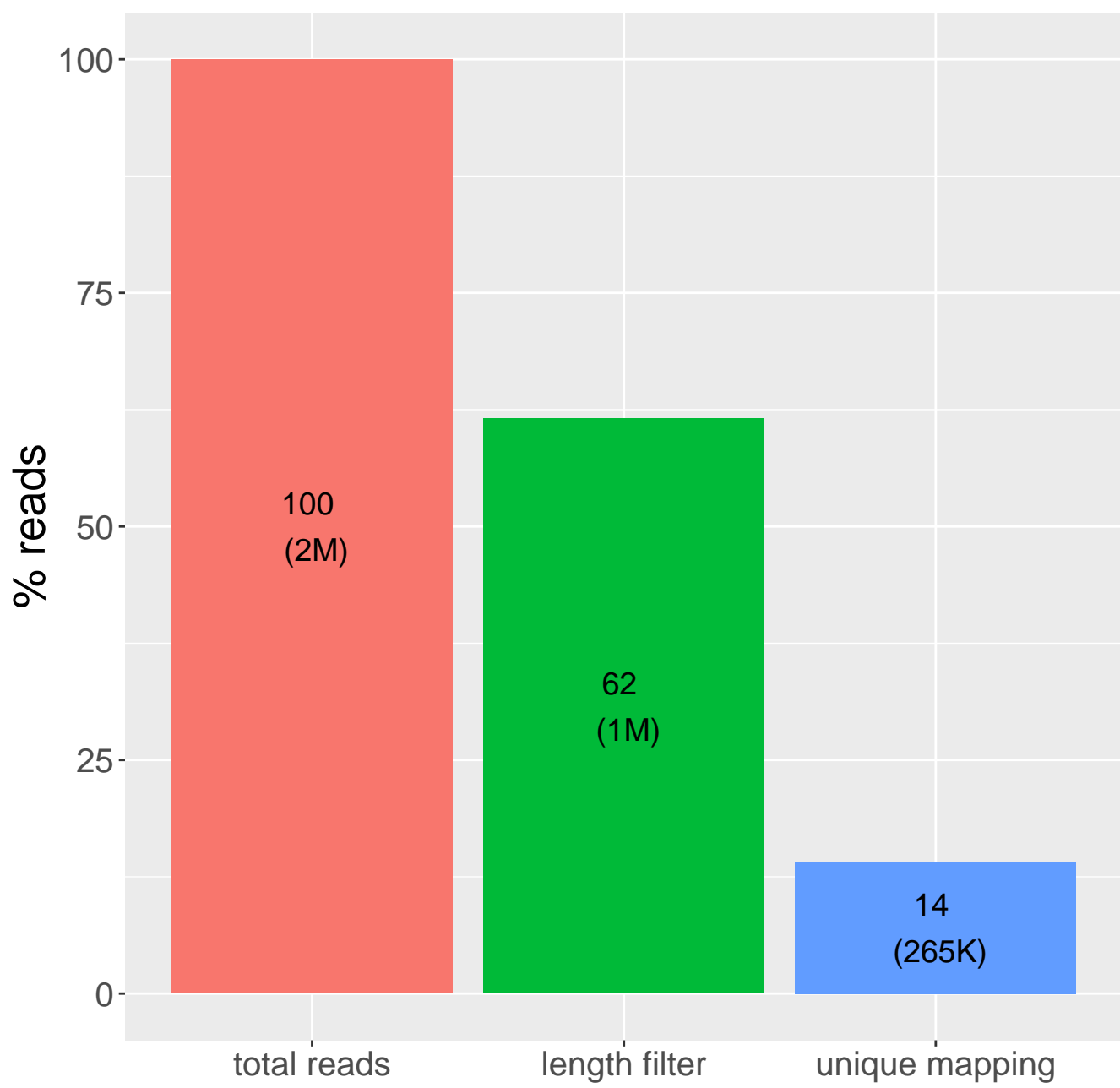
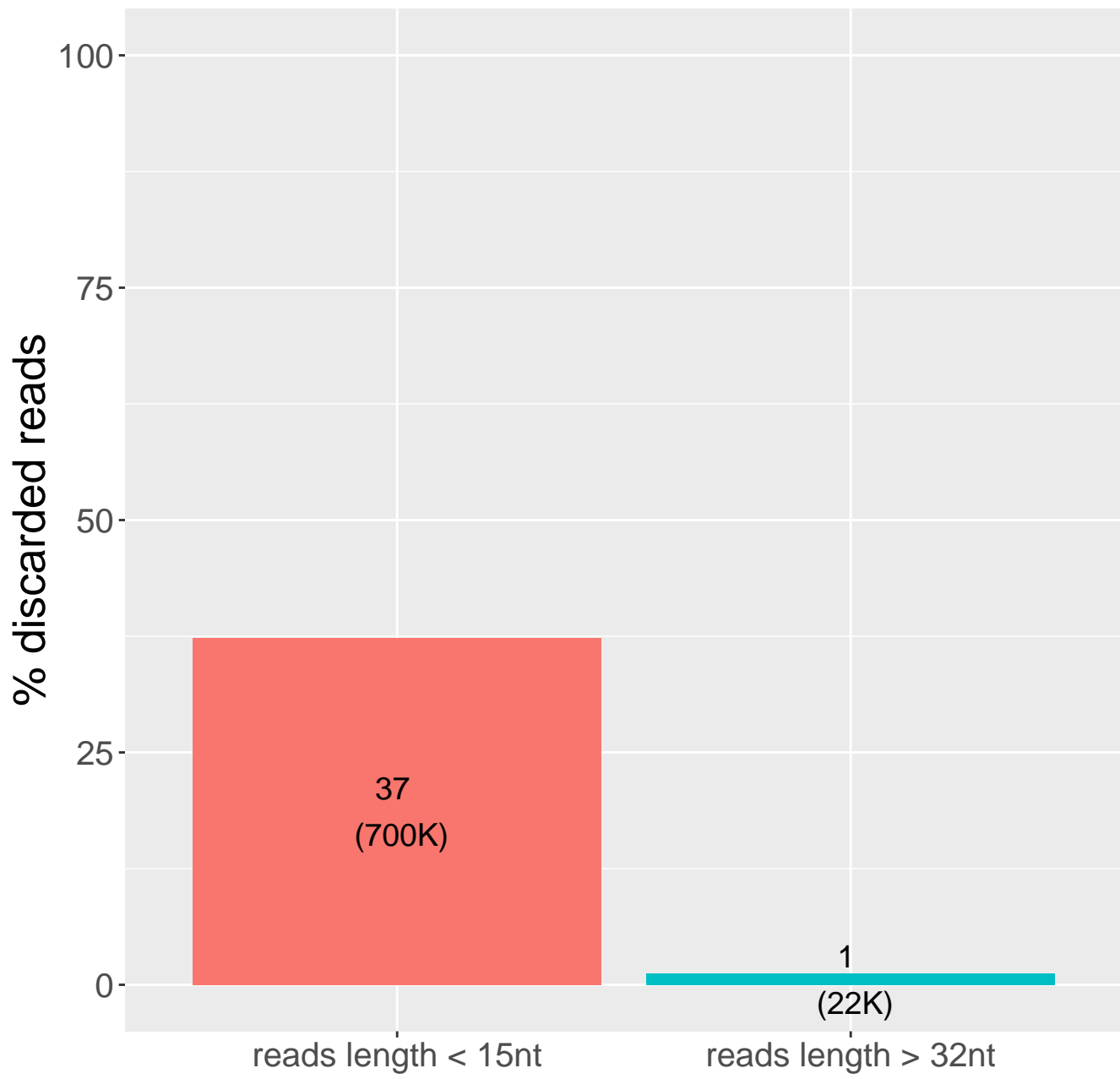


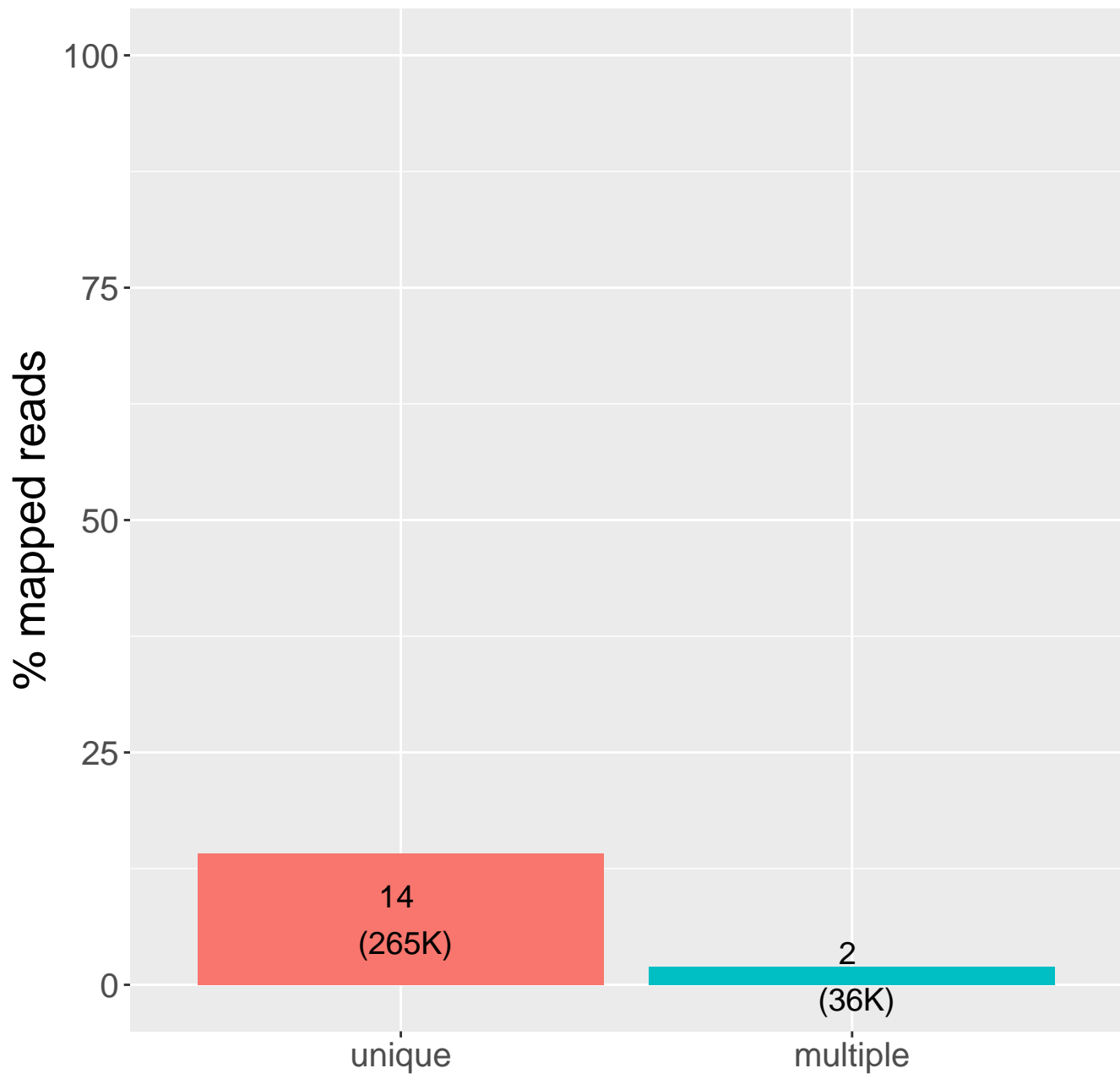
# Retained reads



# Length filtering



# Mapping reads



# Read distribution

% reads

100  
75  
50  
25  
0

miRNA

piRNA

snoRNA

snRNA

rRNA

73  
(193K)

18  
(47K)

3  
(8K)

0  
(1K)

1  
(3K)

