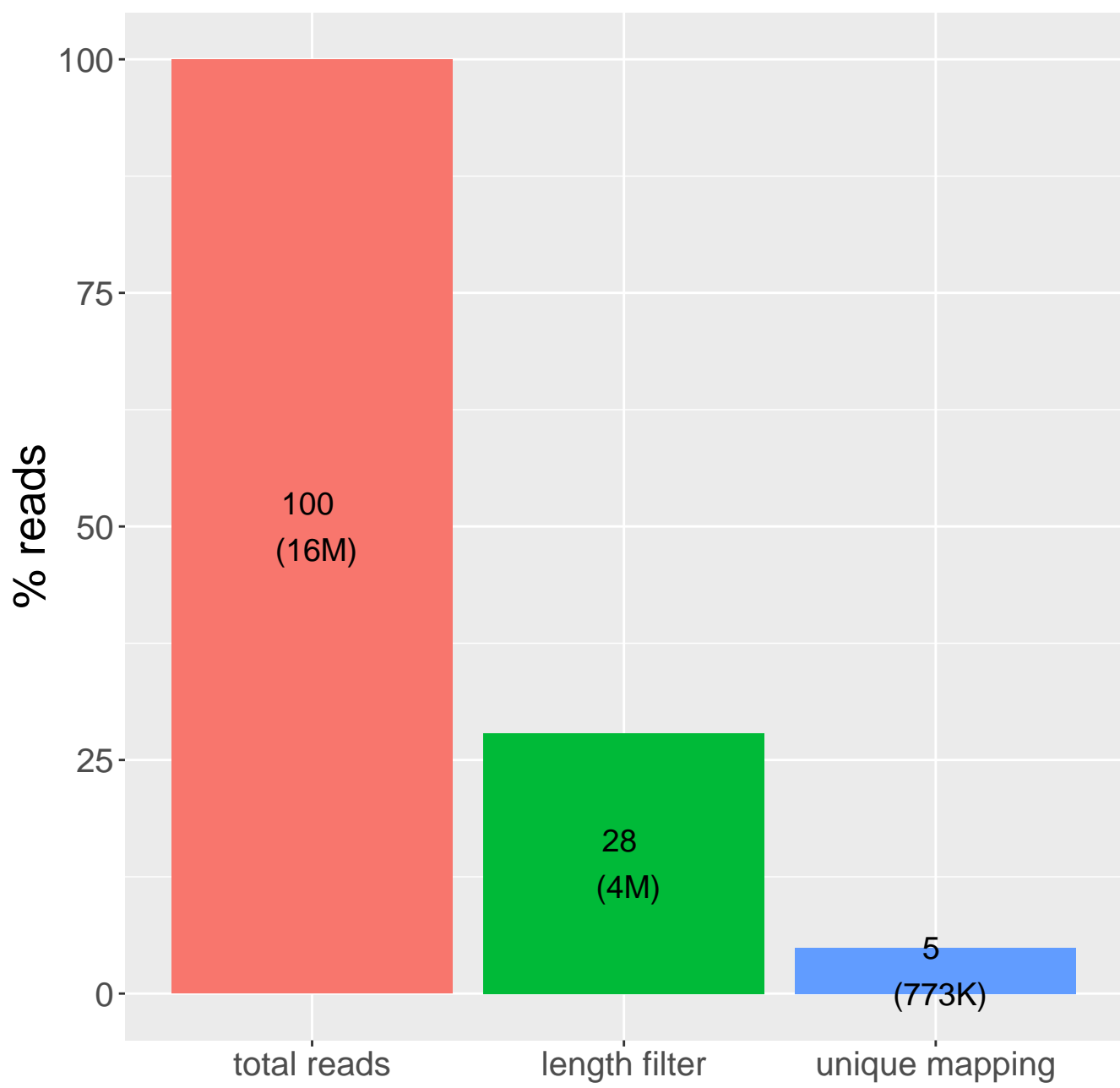
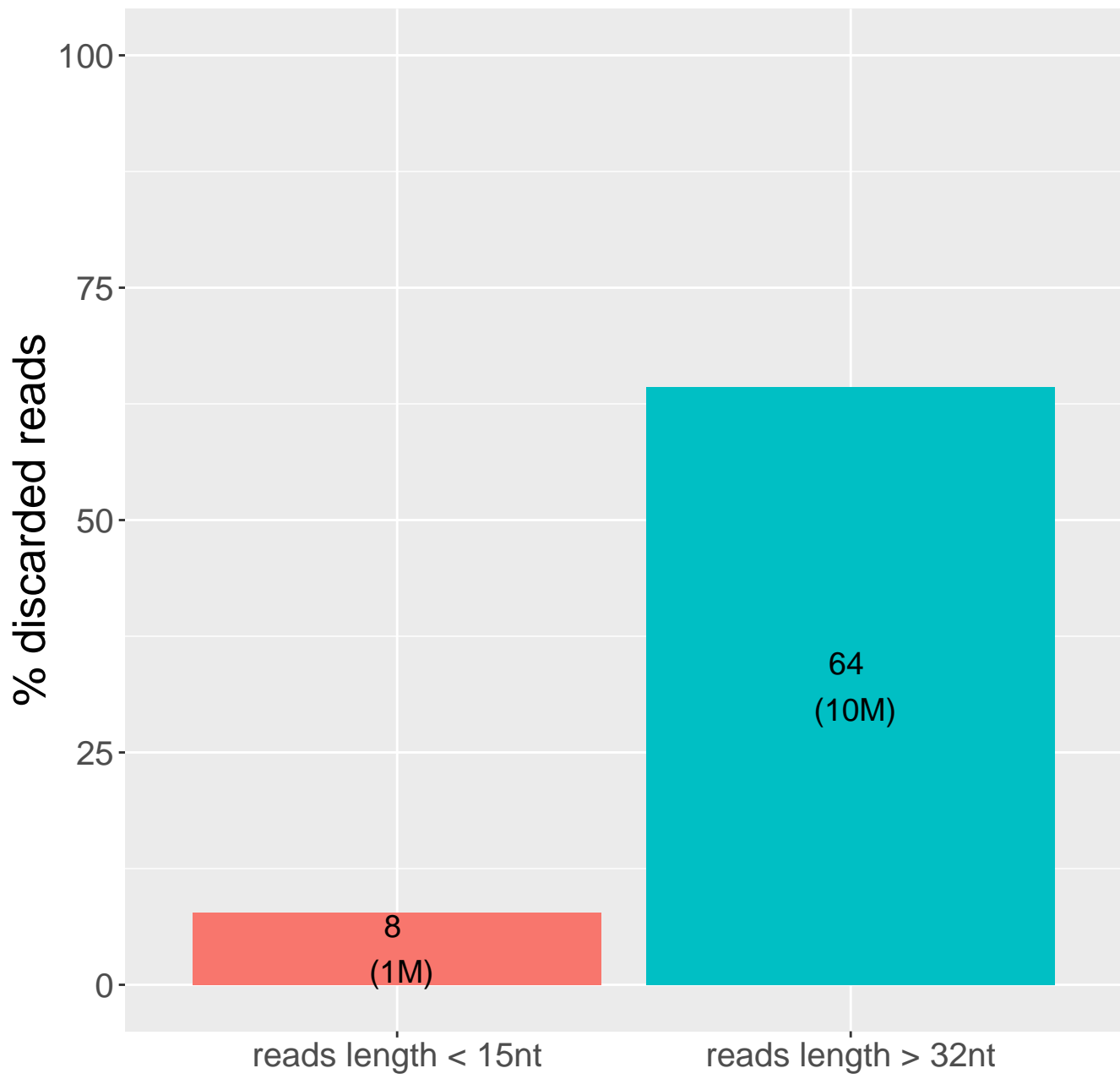


Retained reads

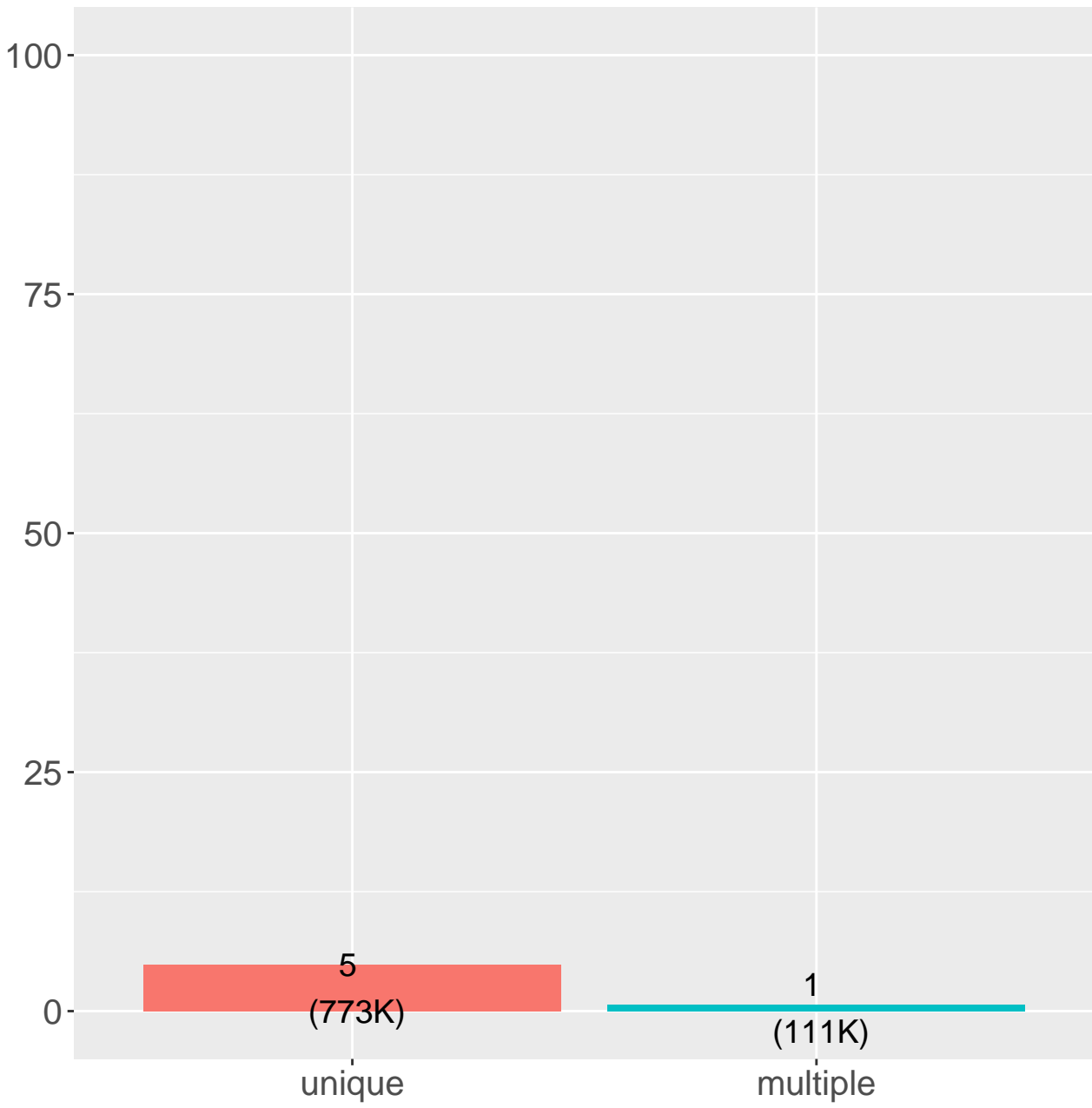


Length filtering



Mapping reads

% mapped reads



Read distribution

% reads

100

75

50

25

0

miRNA

piRNA

snoRNA

snRNA

rRNA

48
(373K)

25
(193K)

3
(21K)

1
(7K)

4
(34K)

