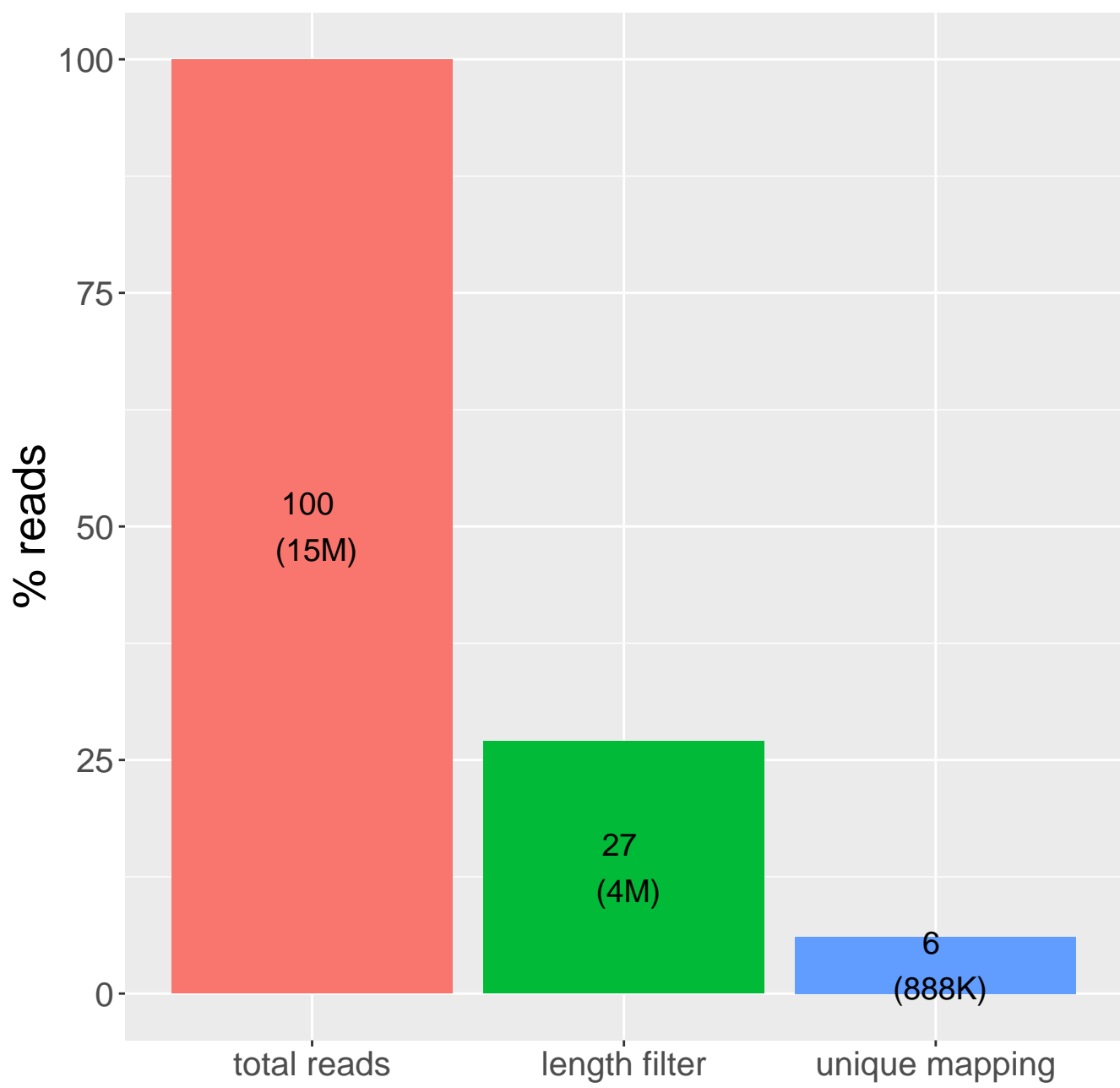
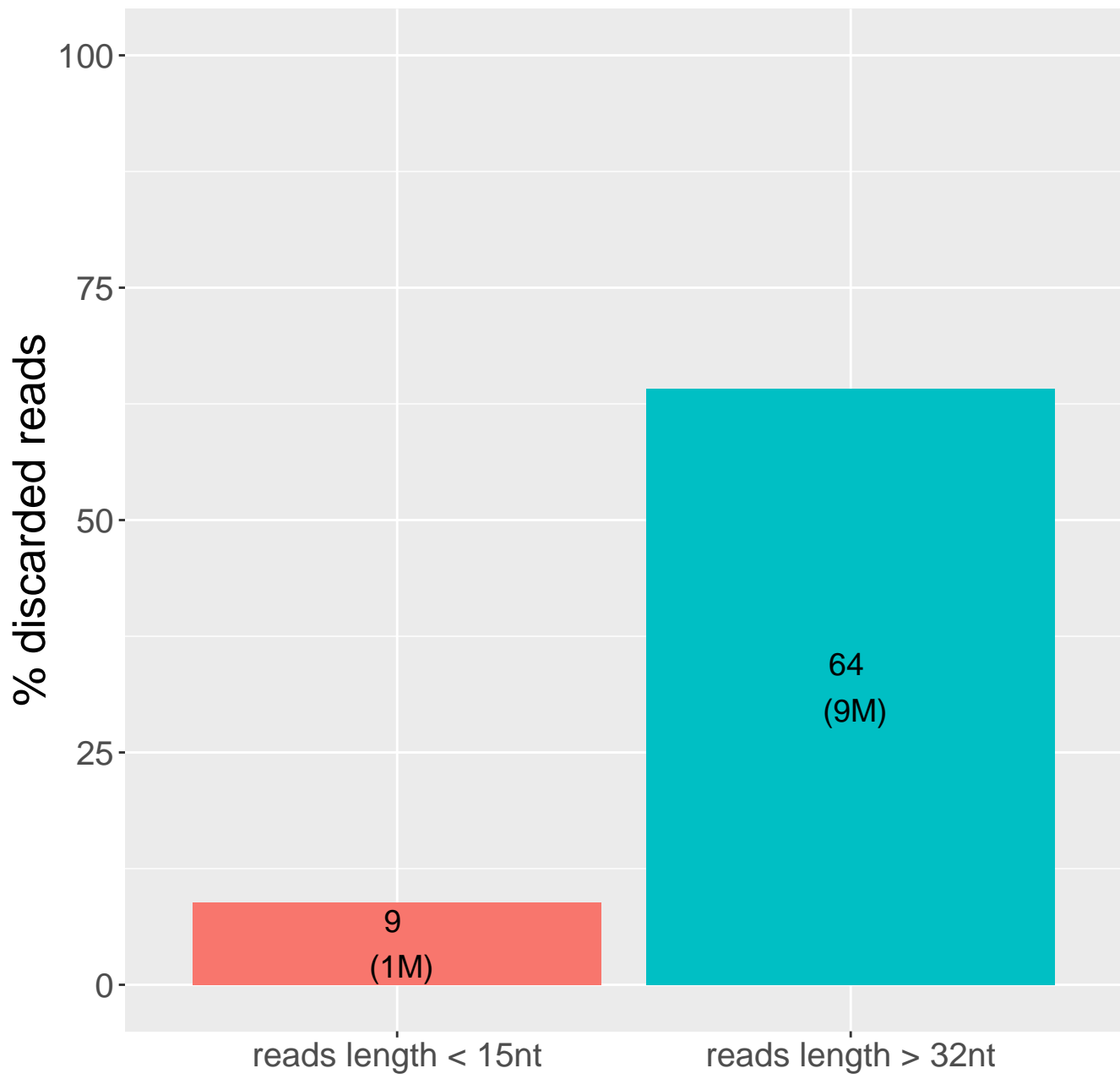


Retained reads

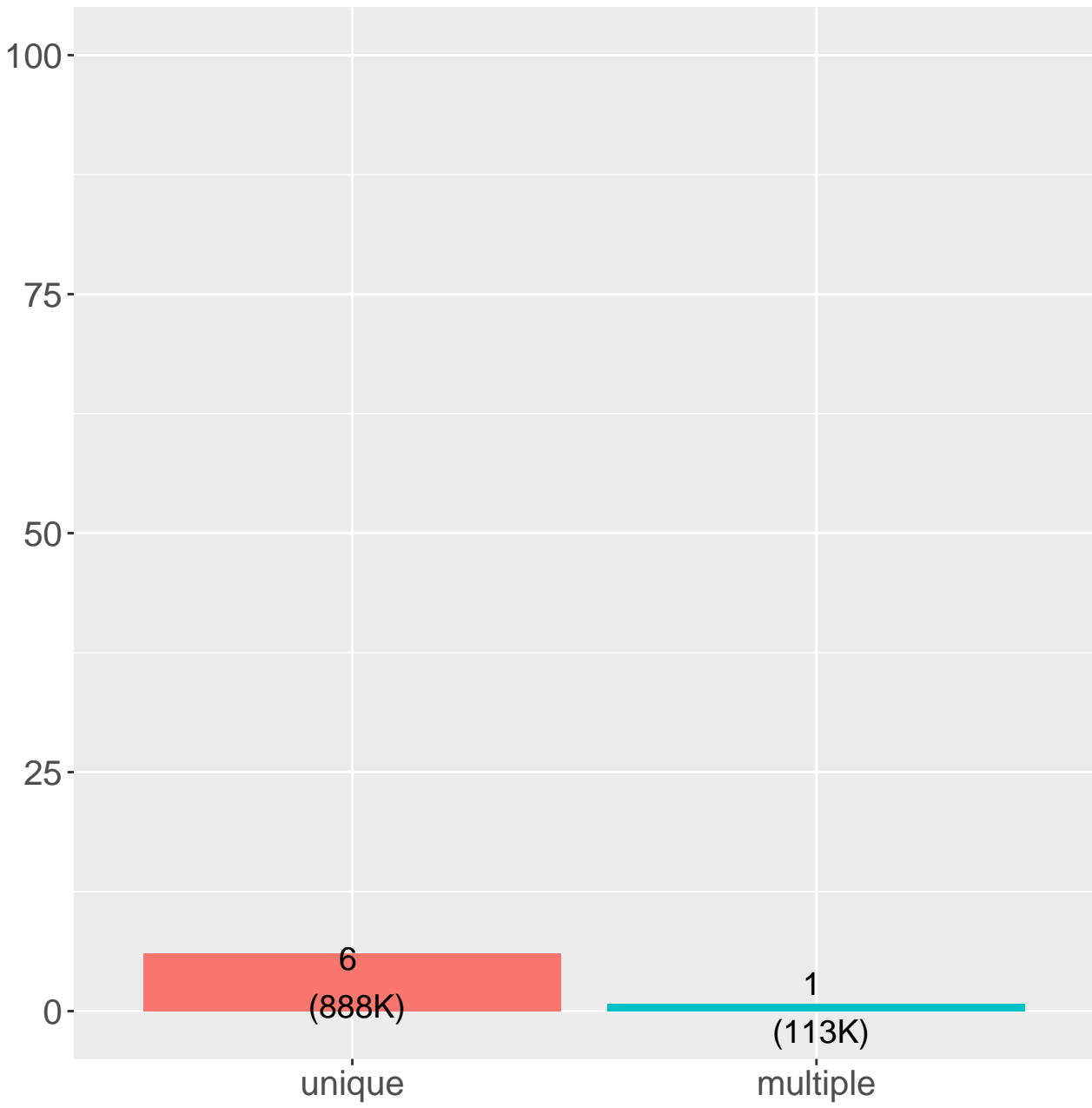


Length filtering



Mapping reads

% mapped reads



Read distribution

% reads

100

75

50

25

0

miRNA

piRNA

snoRNA

snRNA

rRNA

56
(499K)

19
(173K)

4
(37K)

1
(13K)

2
(22K)

