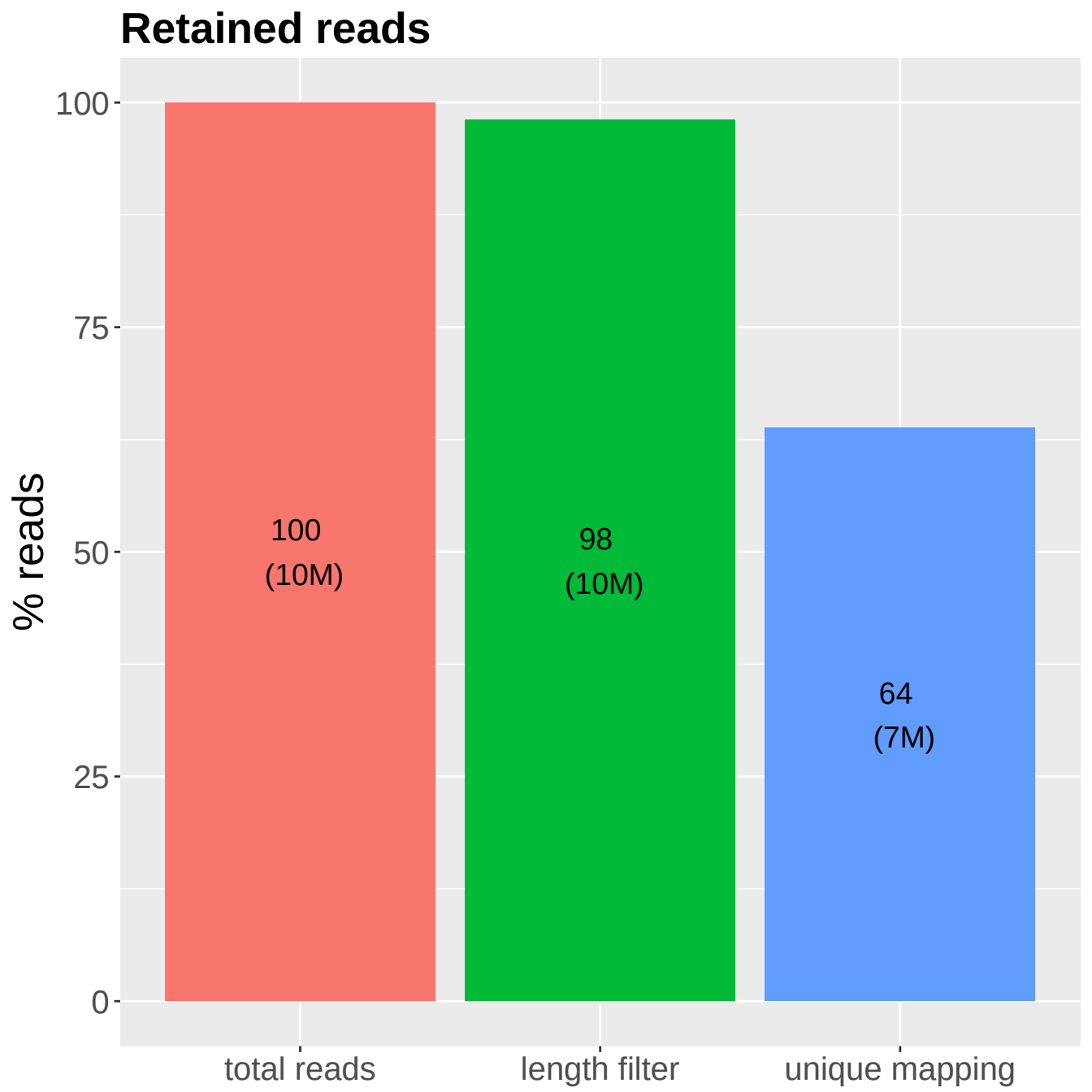
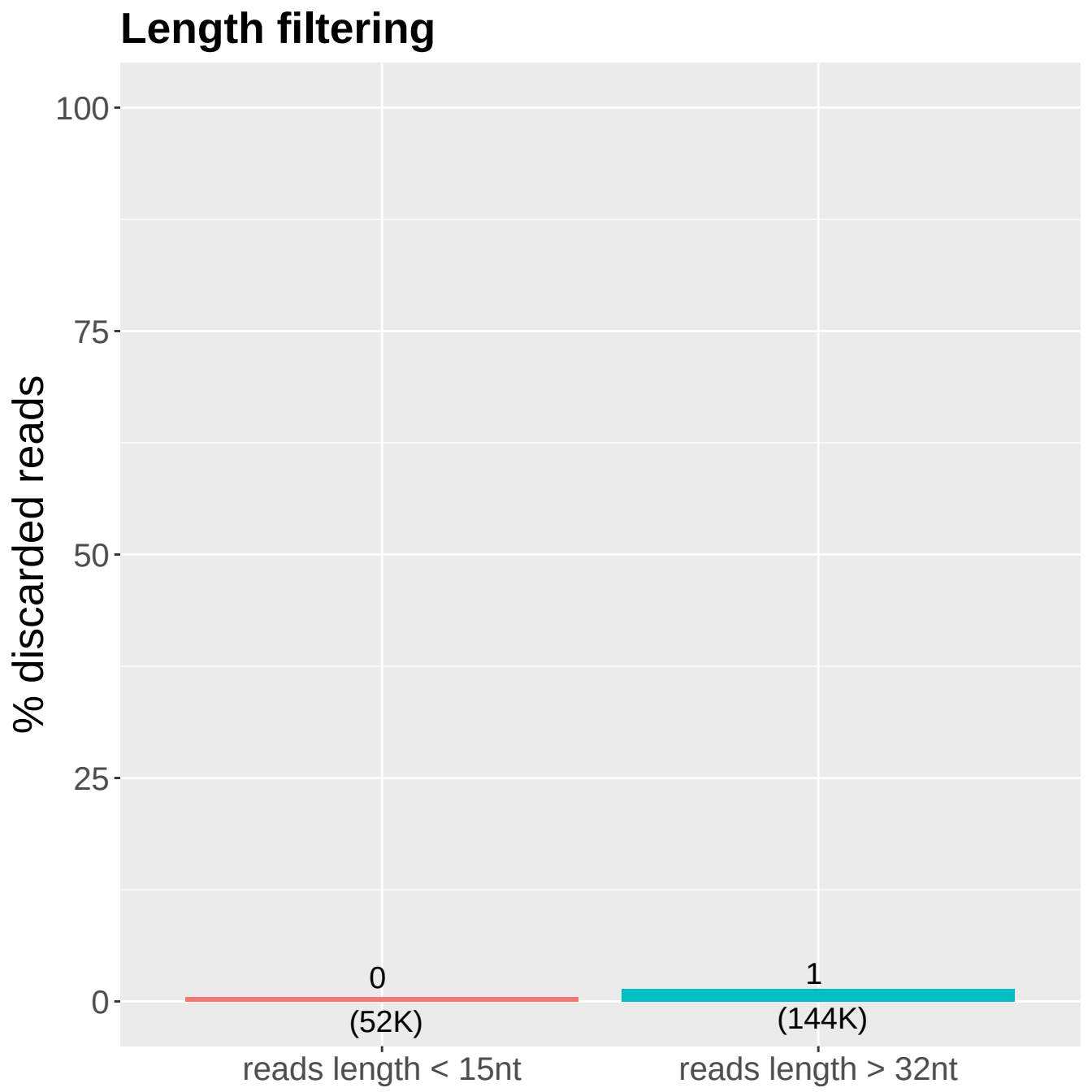
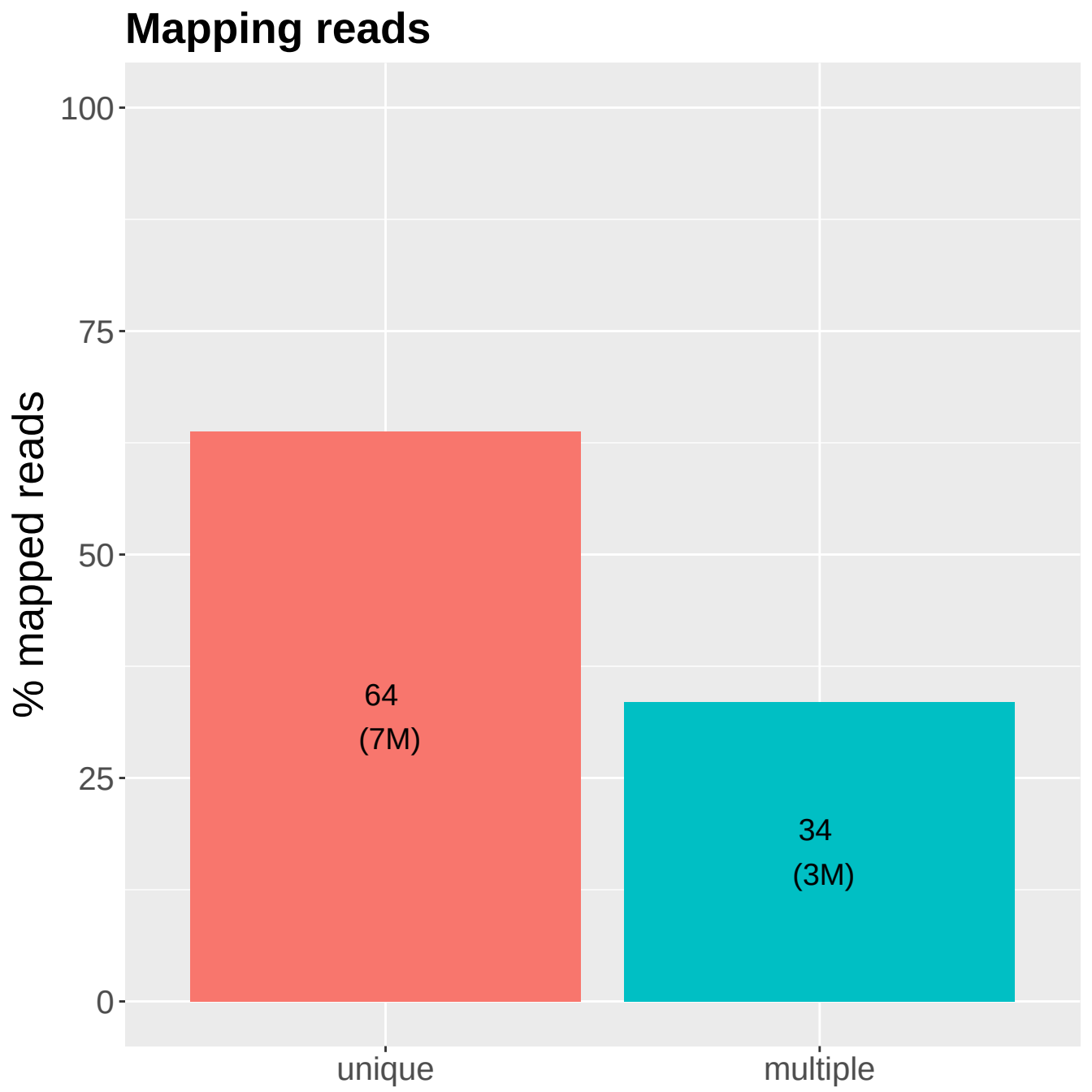
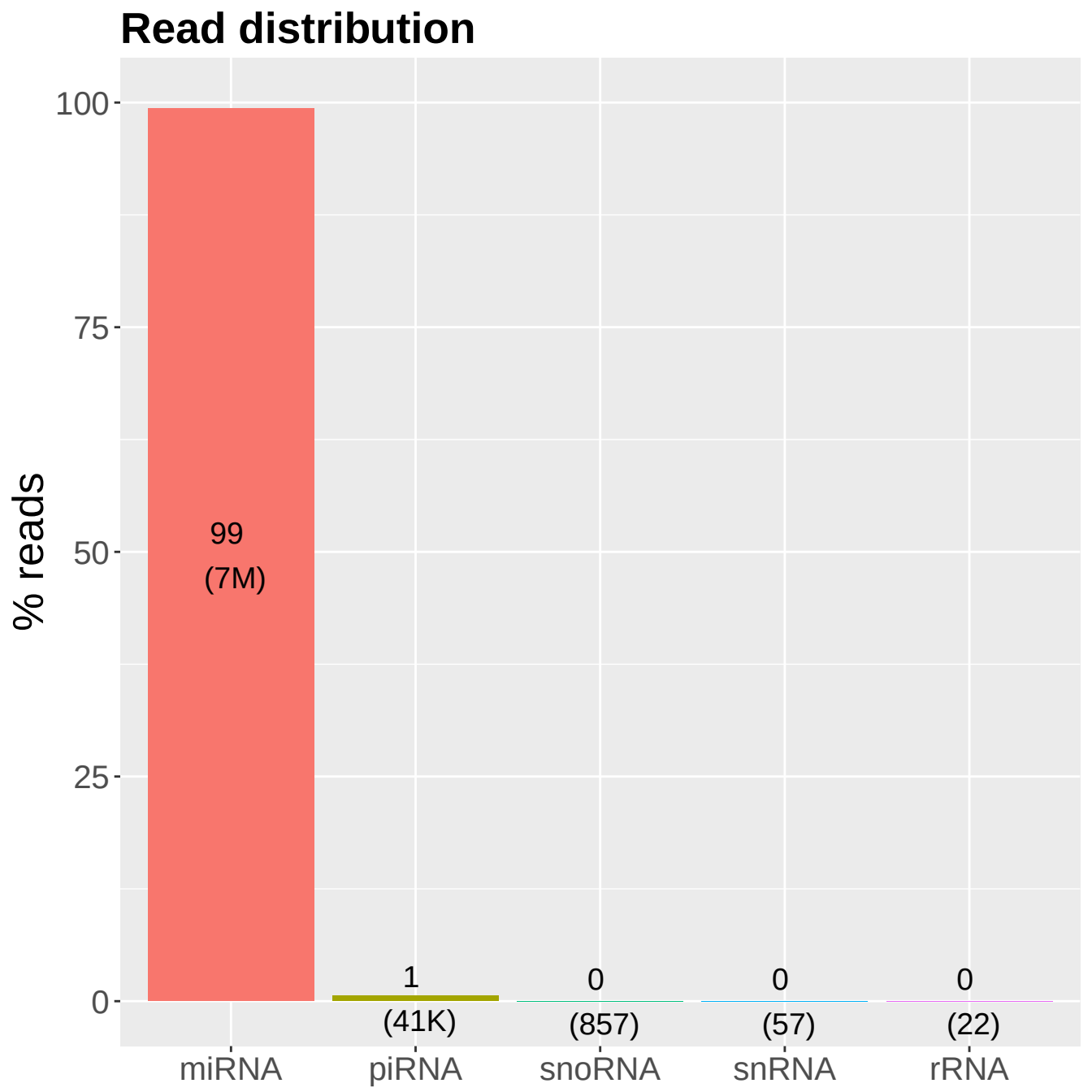

Retained reads
100
75
50
25
0
% reads
100
(10M)
98
(10M)
64
(7M)
total reads
length filter
unique mapping


Length filtering
100
75
50
25
0
% discarded reads
0
(52K)
1
(144K)
reads length < 15nt
reads length > 32nt


Mapping reads
% mapped reads
100
75
50
25
0
64
(7M)
34
(3M)
unique
multiple


Read distribution
100
75
50
25
0
% reads
99
(7M)
1
(41K)
0
(857)
0
(57)
0
(22)
miRNA
piRNA
snoRNA
snRNA
rRNA