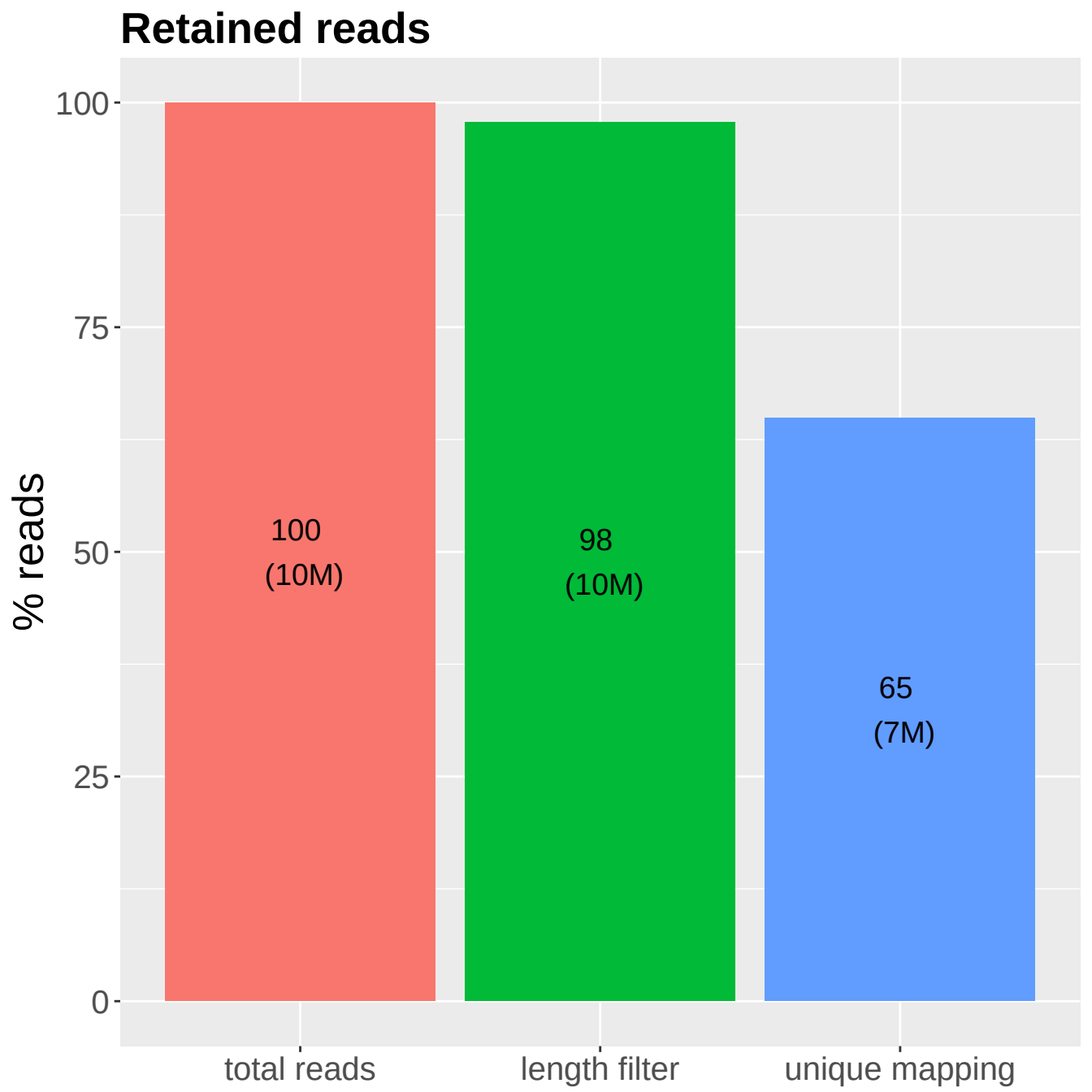
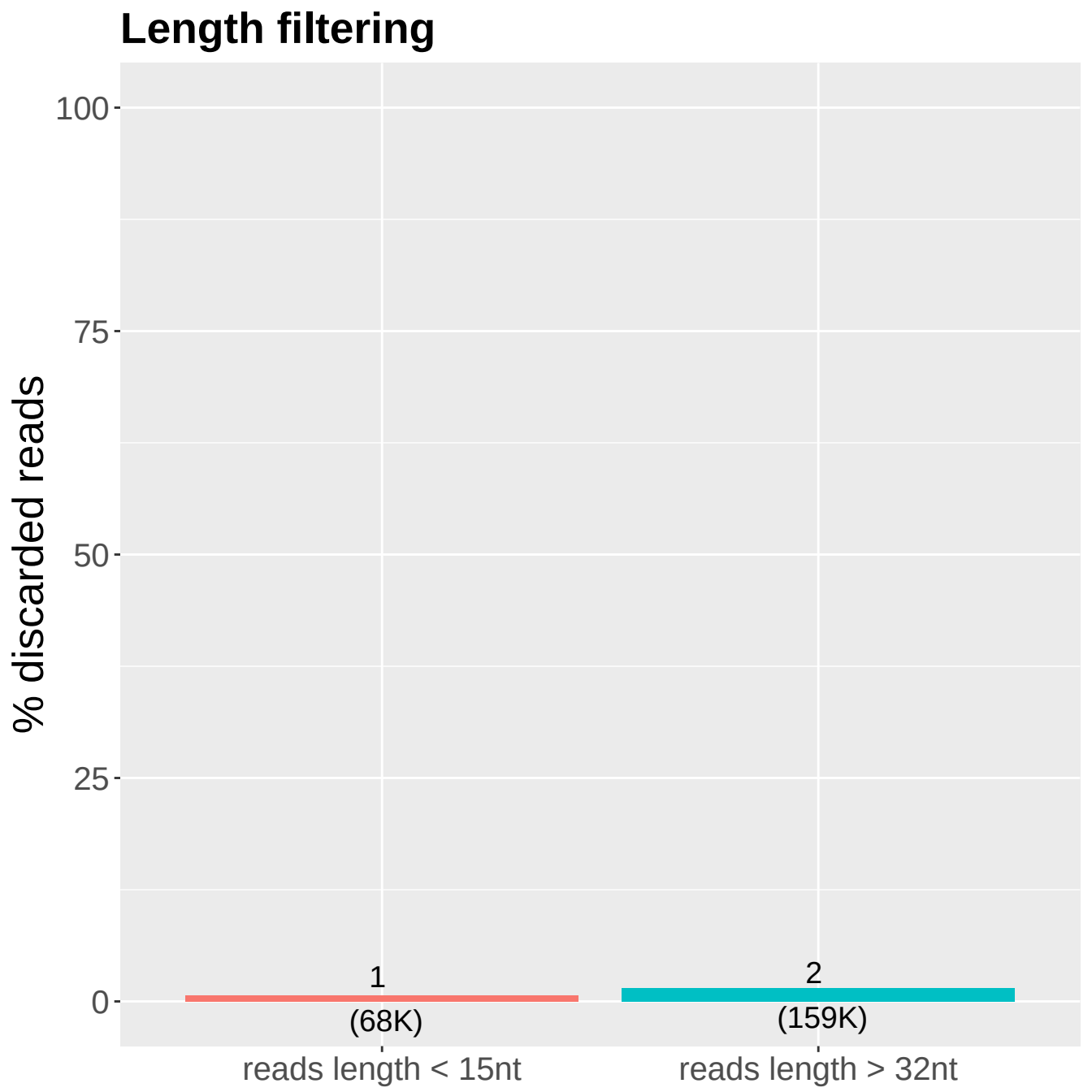
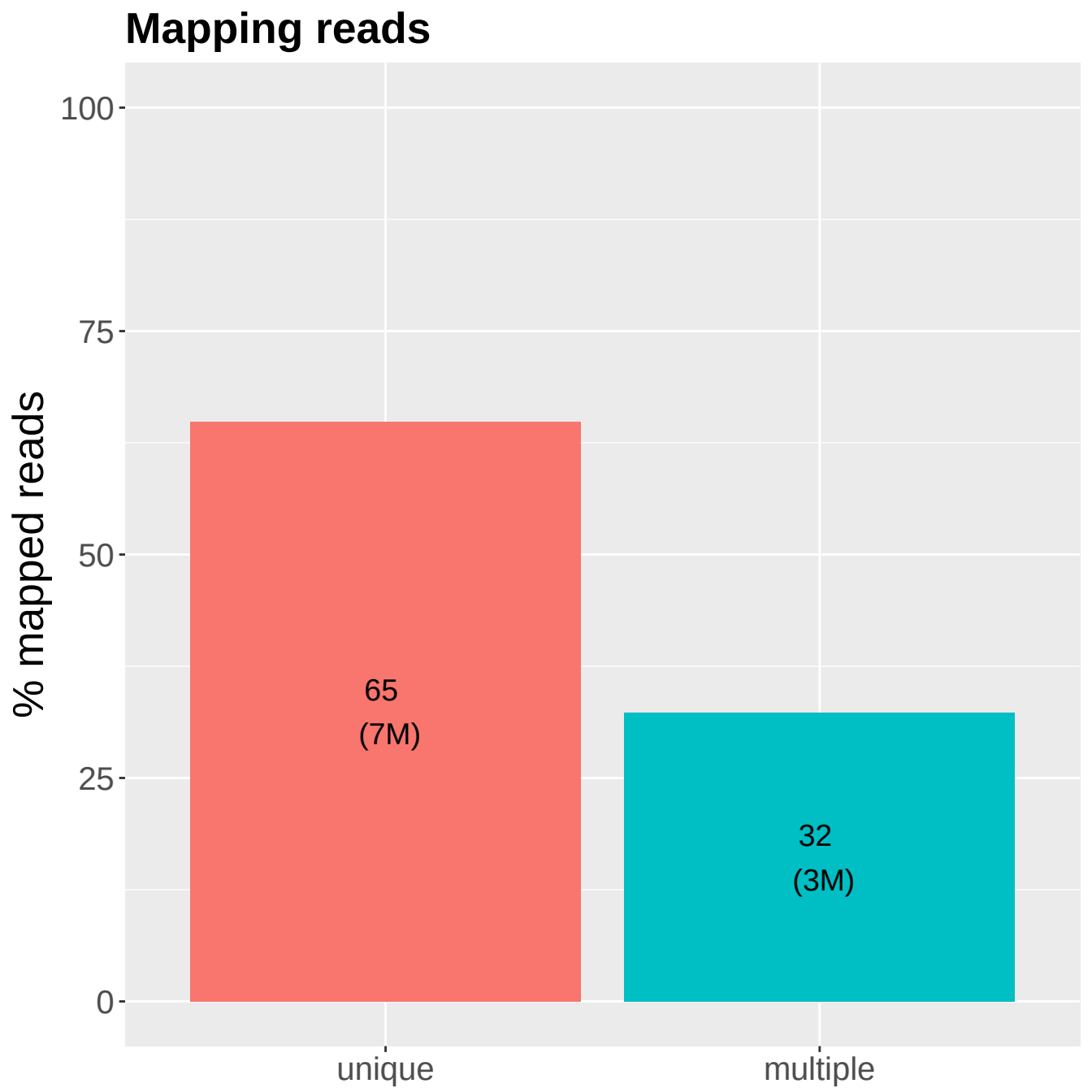
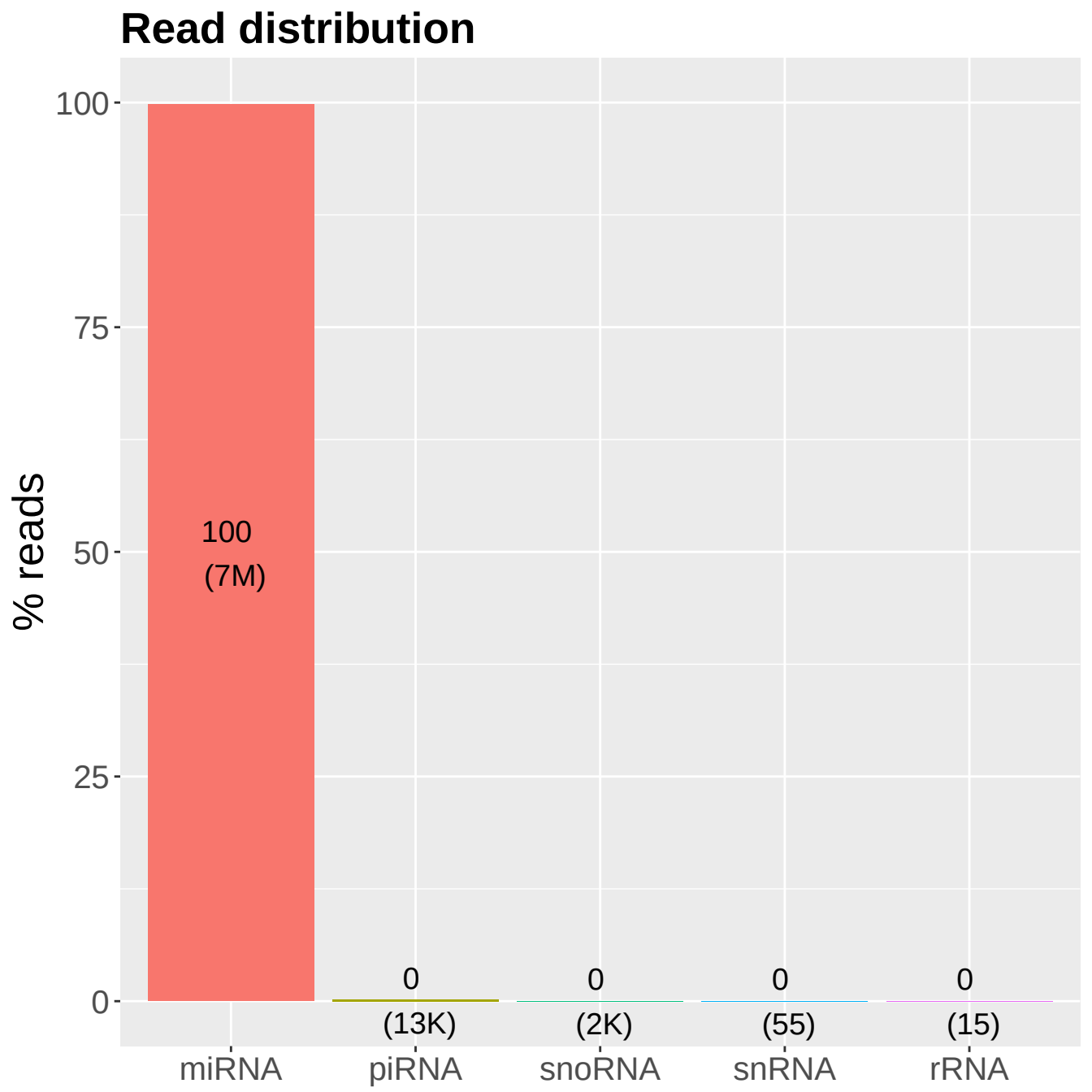

Retained reads
100
75
50
25
0
% reads
100
(10M)
98
(10M)
65
(7M)
total reads
length filter
unique mapping


Length filtering
100
75
50
25
0
% discarded reads
1
(68K)
2
(159K)
reads length < 15nt
reads length > 32nt


Mapping reads
% mapped reads
100
75
50
25
0
65
(7M)
32
(3M)
unique
multiple


Read distribution
100
75
50
25
0
% reads
100
(7M)
0
(13K)
0
(2K)
0
(55)
0
(15)
miRNA
piRNA
snoRNA
snRNA
rRNA